

Mitchell Centre for  
Network Analysis



# An introduction to exponential random graph models (**ERGM**)

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Statistics for Complex and High Dimensional Systems, Eindhoven

ERGM: probability model for **adjacency matrices**  
with pmf:

$$p(x) = \exp\{\theta^T z(x) - \psi(\theta)\}$$

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Normalising constant:  $\psi(\theta)$

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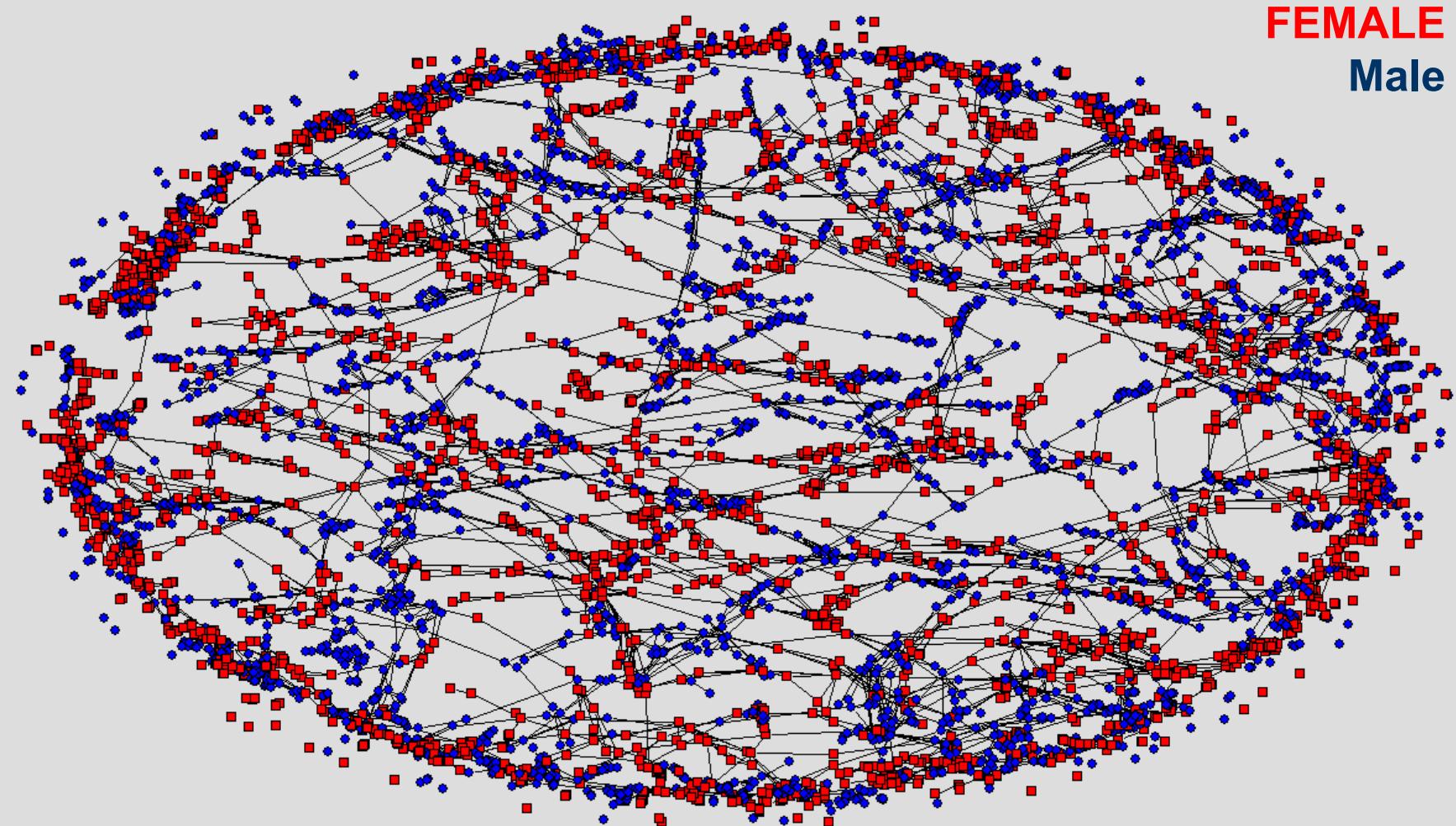
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... it is an exponential family distribution (hence ERGM)

## Part 1a

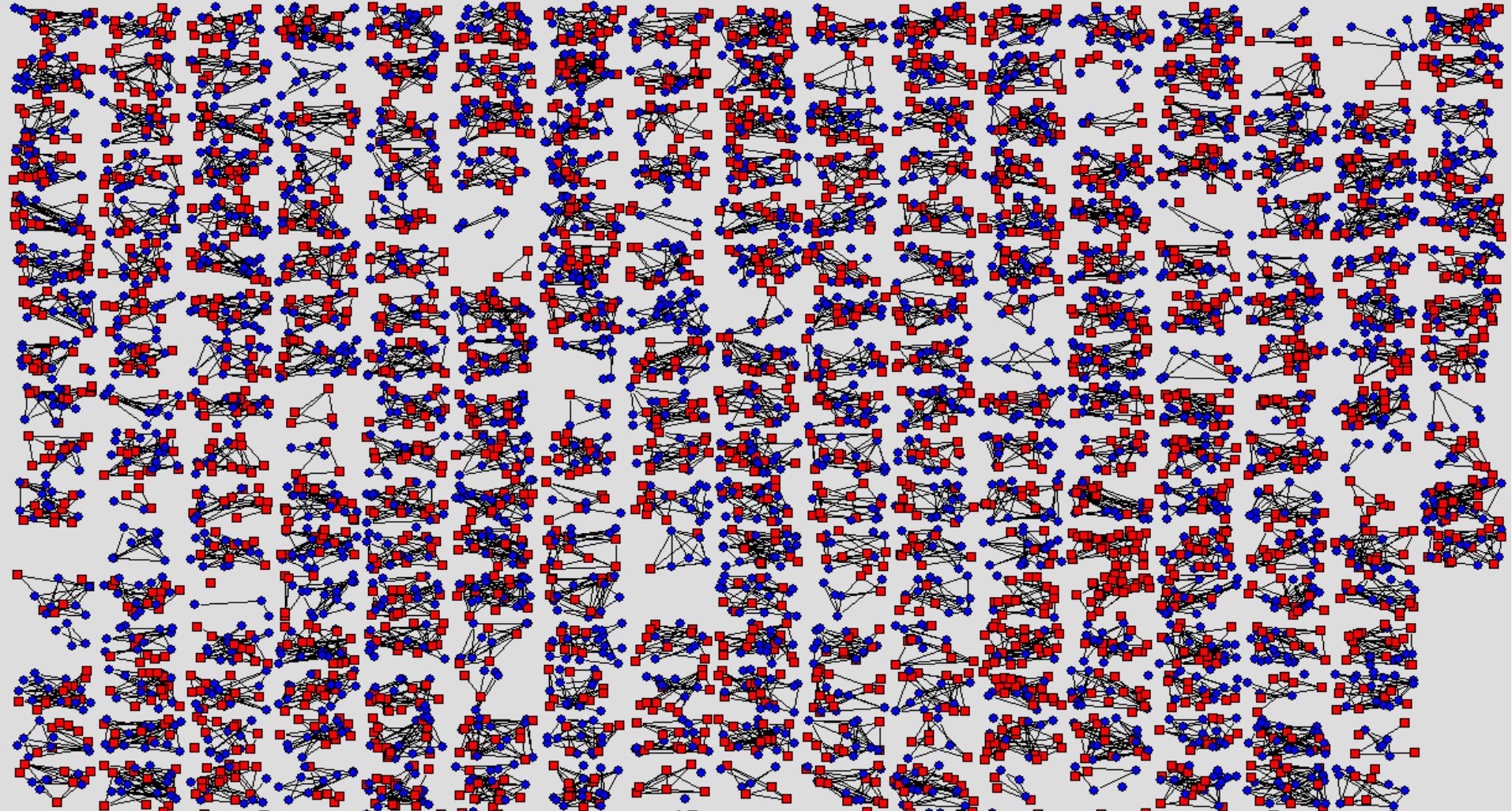
### Why an ERGM

# Networks matter – ERGMS matter



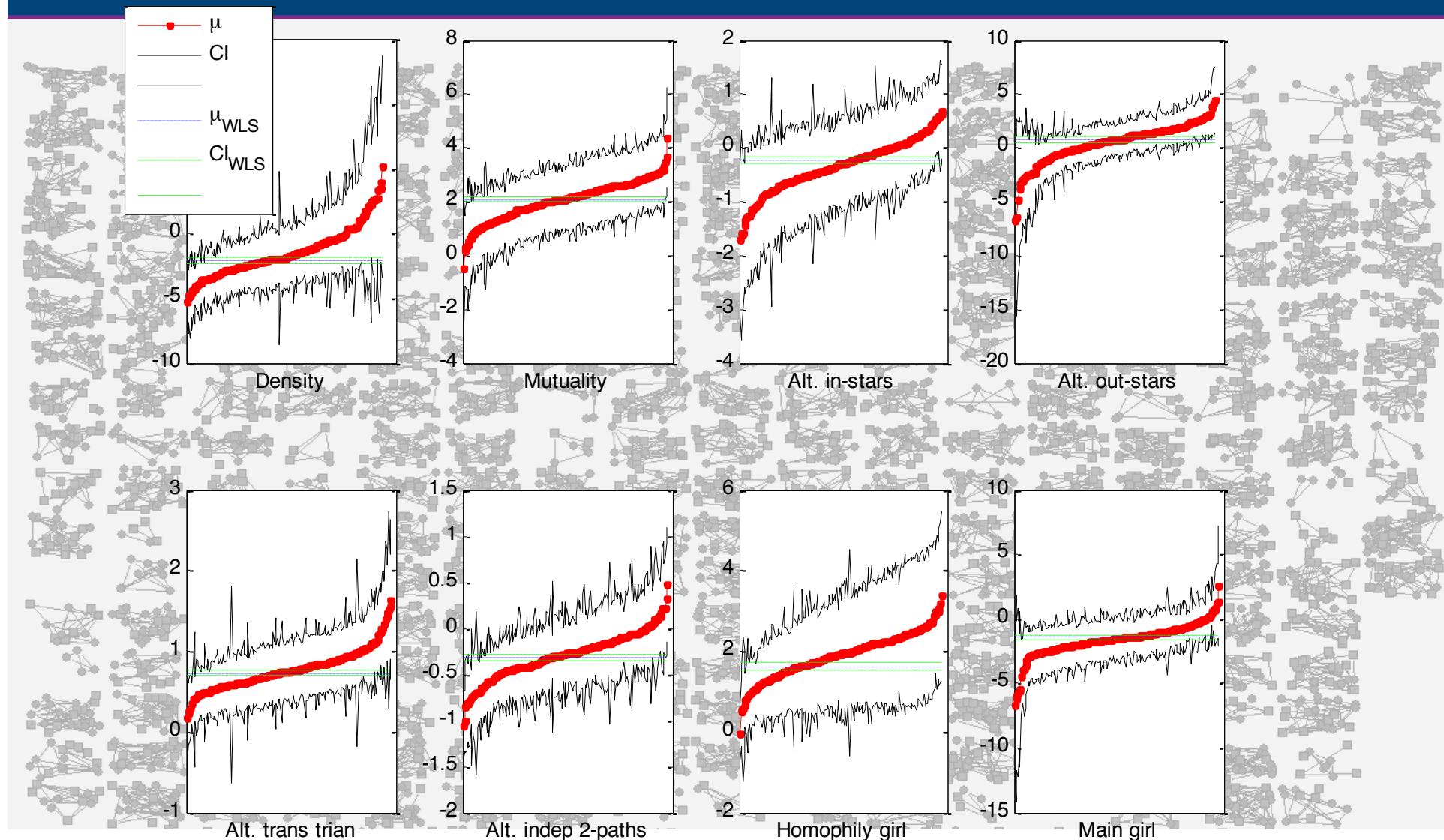
**6018 grade 6 children 1966**

# Networks matter – ERGMS matter



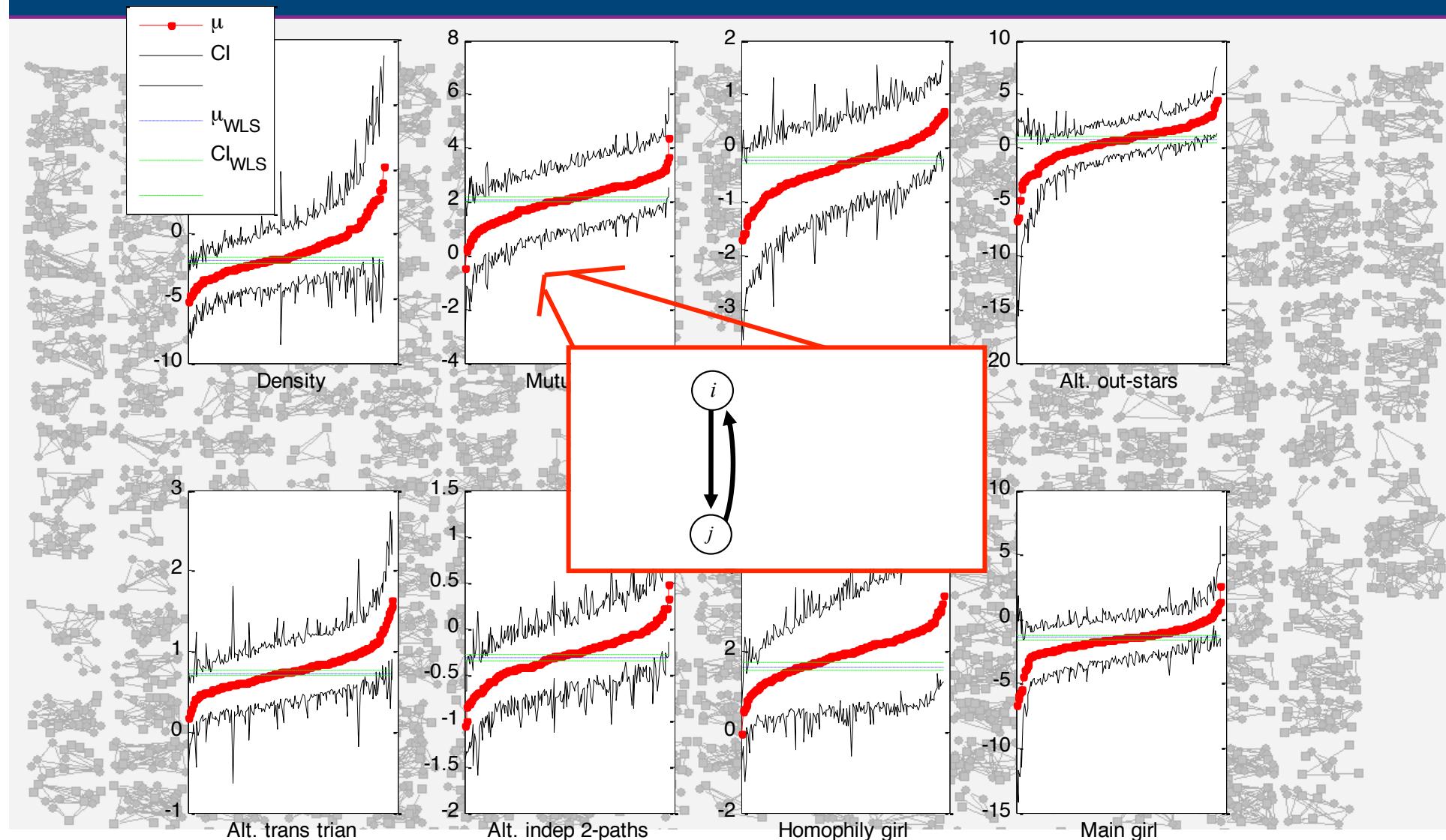
6018 grade 6 children 1966 – 300 schools Stockholm

# Networks matter – ERGMS matter



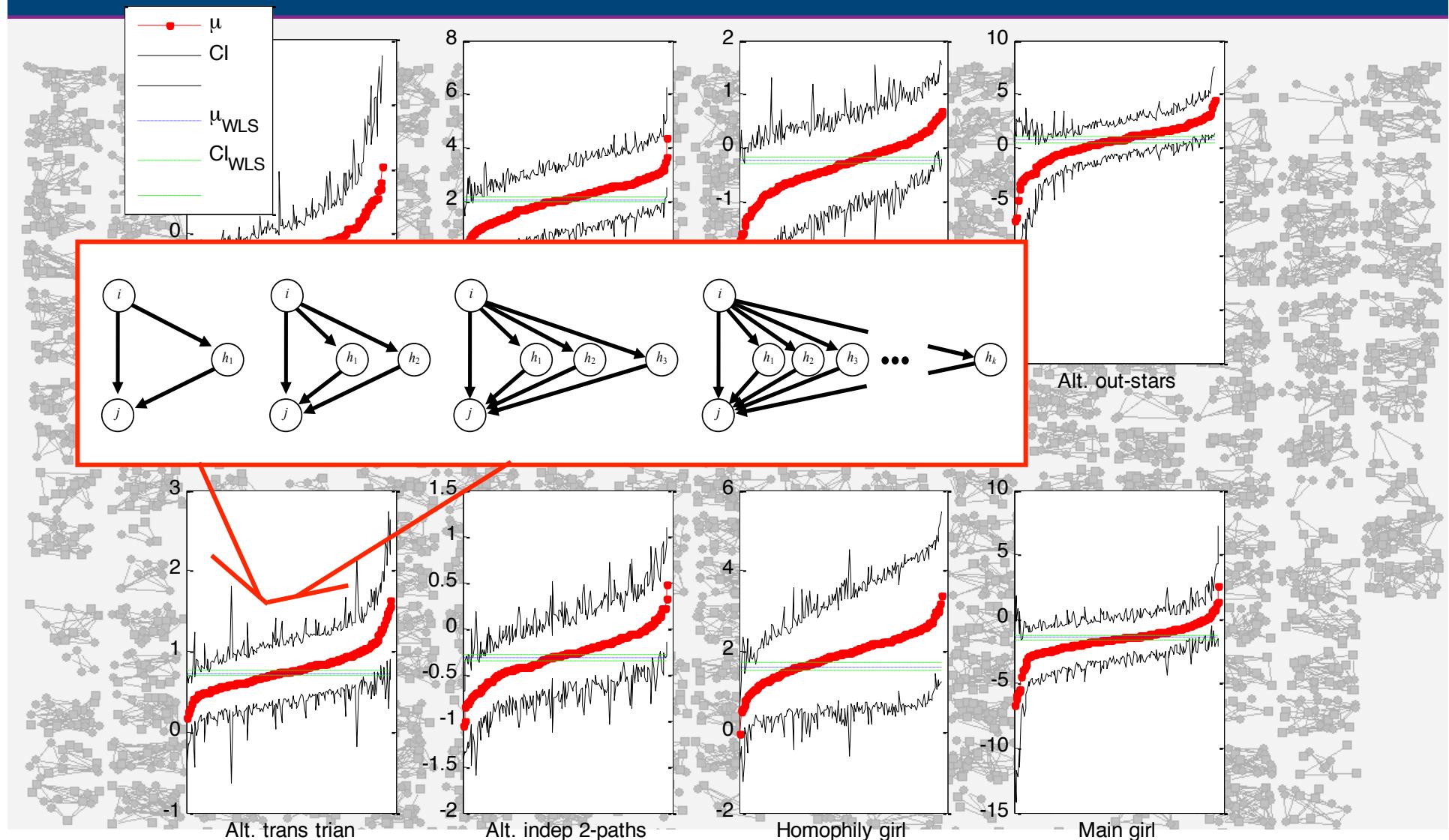
**6018 grade 6 children 1966 – 200 schools Stockholm**  
*Koskinen and Stenberg (in press) JEBS*

# Networks matter – ERGMS matter



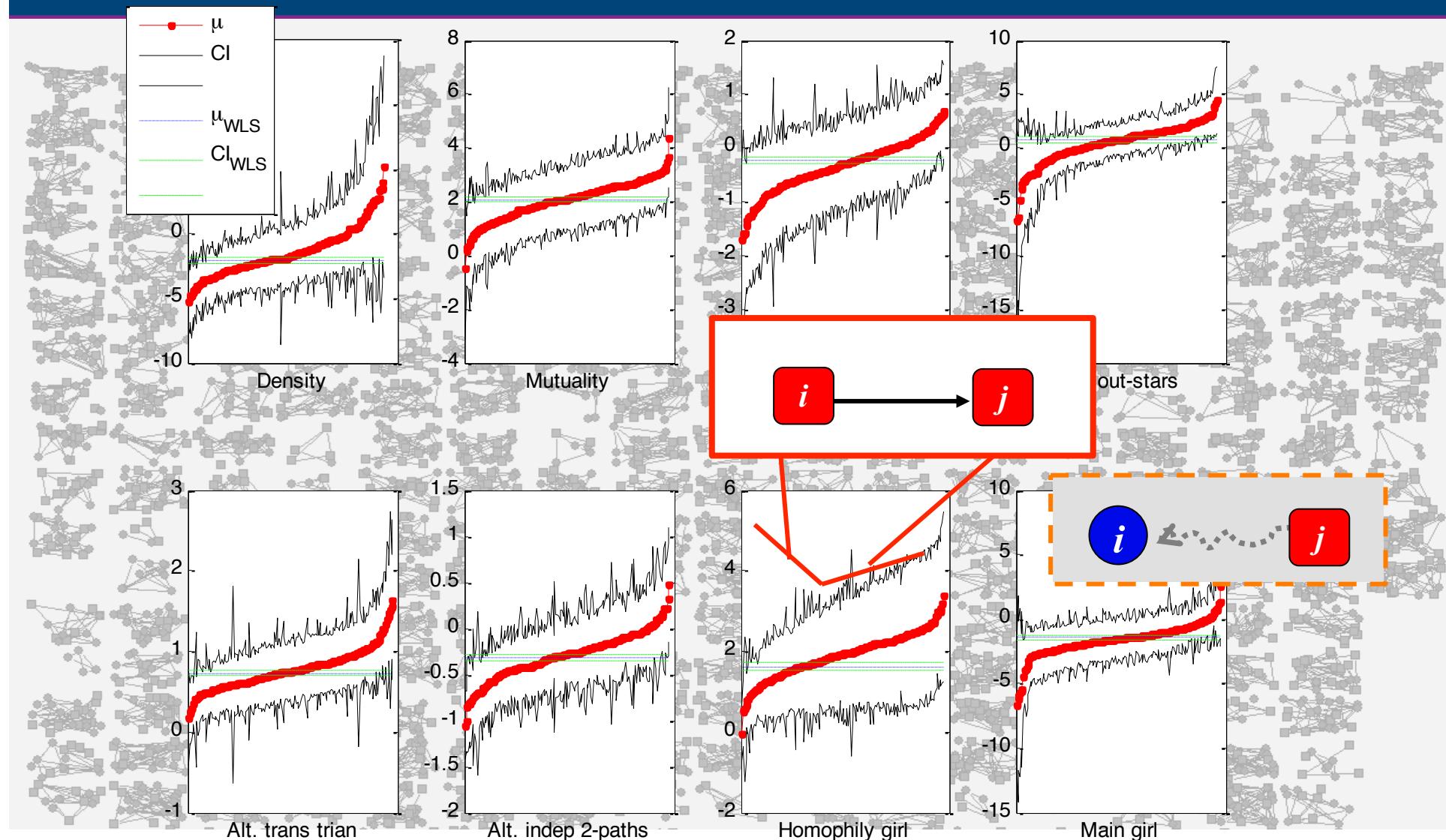
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# Networks matter – ERGMS matter



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## Part 1b

### Minimum learning outcomes

- Get a working handle on what we are trying to model
- Familiarise ourselves with common model specifications
- Fit our first models in statnet and PNet
- Fitting procedure
  - Estimation
  - Convergence check
  - Goodness of fit
- Orientation about future developments

# Part 1c

## Modelling graphs

- Notational preliminaries
- Why and what is an ERGM
- Dependencies
- Estimation
  - Geyer-Thompson
  - Robins-Monro
  - Bayes (The issue, Moller et al, LISA, exchange algorithm)
- Interpretation of effects
- Convergence and GOF
- Further issues

Numerous recent substantively driven studies

- Gondal, The local and global structure of knowledge production in an emergent research field, **SOCNET 2011**
- Lomi and Palotti, Relational collaboration among spatial multipoint competitors , **SOCNET 2011**
- Wimmer & Lewis, Beyond and Below Racial Homophily, **AJS 2010**
- Lusher, Masculinity, educational achievement and social status, **GENDER & EDUCATION 2011**
- Rank et al. (2010). Structural logic of intra-organizational networks, **ORG SCI, 2010.**

**Book for applied researchers:** Lusher, Koskinen, Robins  
ERGMs for SN, CUP, 2011

Exponential random graph models (ERGMs) are increasingly applied to observed network data and are central to understanding social structure and network processes. The chapters in this edited volume provide the theoretical and methodological underpinnings of ERGMs, including models for univariate, multivariate, bipartite, longitudinal, and social-influence type ERGMs. Each method is applied in individual case studies illustrating how social science theories may be examined empirically using ERGMs. The authors supply the reader with sufficient detail to specify ERGMs, fit them to data with any of the available software packages, and interpret the results.

Dr. Dean Lusher is Lecturer in Sociology at Swinburne University of Technology. He works closely with leading methodologists to develop an intuitive understanding of exponential graph models, how they link to broader network theory, and how to fit them to real-life data. His research applications are directed at issues of social norms and social hierarchies.

Dr. Johan Koskinen is Lecturer in Social Sciences at the University of Manchester. He is a statistician working with statistical modeling and inference. Focusing on social network data, Dr. Koskinen deals with generative models for different types of structures, such as longitudinal network data, networks nested in multilevel structures, and multilevel networks classified by affiliations.

Garry Robins is Professor in the School of Psychological Sciences at the University of Melbourne. Robins is a mathematical psychologist whose research deals with quantitative and statistical models for social and relational systems. His research has won international awards from the Psychometric Society, the American Psychological Association, and the International Network for Social Network Analysis.

# Exponential Random Graph Models for Social Networks

Lusher, Koskinen  
Robins  
**Exponential Random Graph Models for Social Networks**

THEORIES, METHODS, AND APPLICATIONS

Dean Lusher, Johan Koskinen,  
Garry Robins

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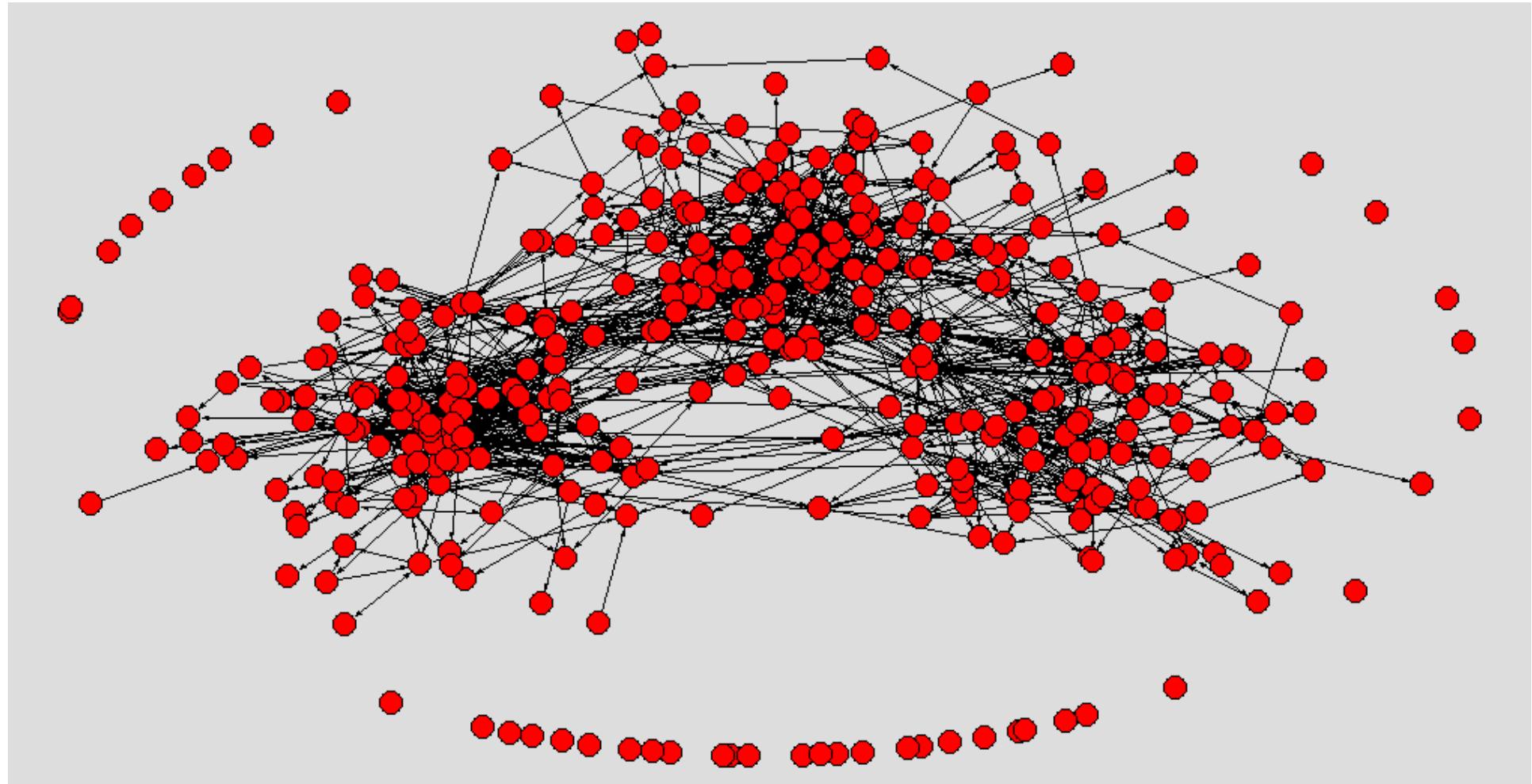
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# The exponential random graph model ( $p^*$ ) framework

- An ERGM ( $p^*$ ) model is a statistical model for the **ties** in a network
- Independent (pairs of) ties (p1, Holland and Leinhardt, 1981; Fienberg and Wasserman, 1979, 1981)
- Markov graphs (Frank and Strauss, 1986)
- Extensions (Pattison & Wasserman, 1999; Robins, Pattison & Wasserman, 1999; Wasserman & Pattison, 1996)
- New specifications (Snijders et al., 2006; Hunter & Handcock, 2006)

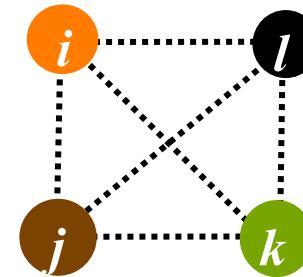
# ERGMS – modelling graphs



- We want to model **tie variables**
- But **structure** – overall pattern – is evident
- What kind of **structural elements** can we incorporate in the model for the tie variables?

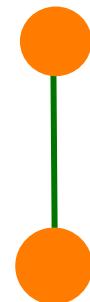
# ERGMS – modelling graphs

If we believe that the frequency of interaction/**density** is an important aspect of the network



We should include

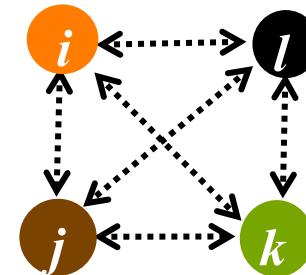
Counts of



the number of ties in our model

# ERGMS – modelling graphs

If we believe that the **reciprocity** is an important aspect of the (directed) network



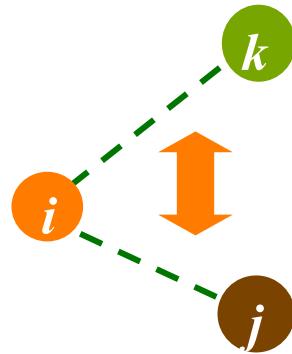
We should include

Counts of



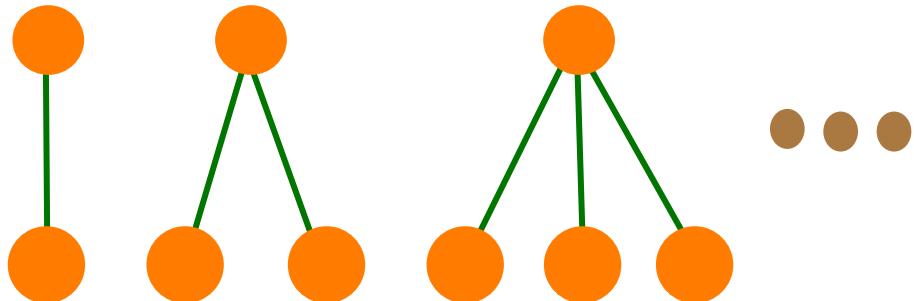
the number of mutual ties in our model

If we believe that an important aspect of the network is that

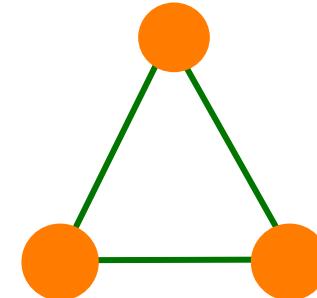


two edge indicators  $\{i,j\}$  and  $\{i',k\}$  are conditionally **dependent** if  $\{i,j\} \cap \{i',k\} \neq \emptyset$

We should include counts of



*degree distribution; preferential attachment,  
etc*

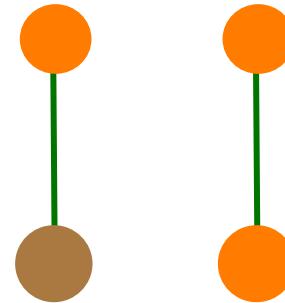


*friends meet through friends; clustering; etc*

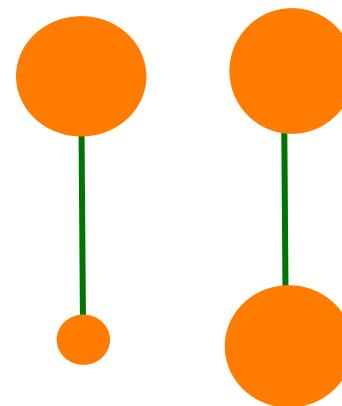
## ERGMS – modelling graphs

If we believe that the **attributes** of the actors are important (selection effects, homophily, etc)

We should include counts of



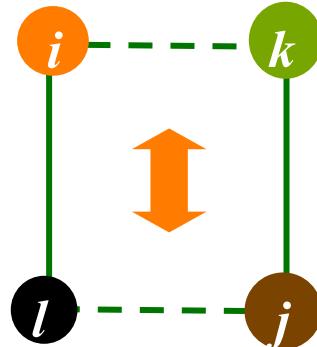
*Heterophily/homophily*



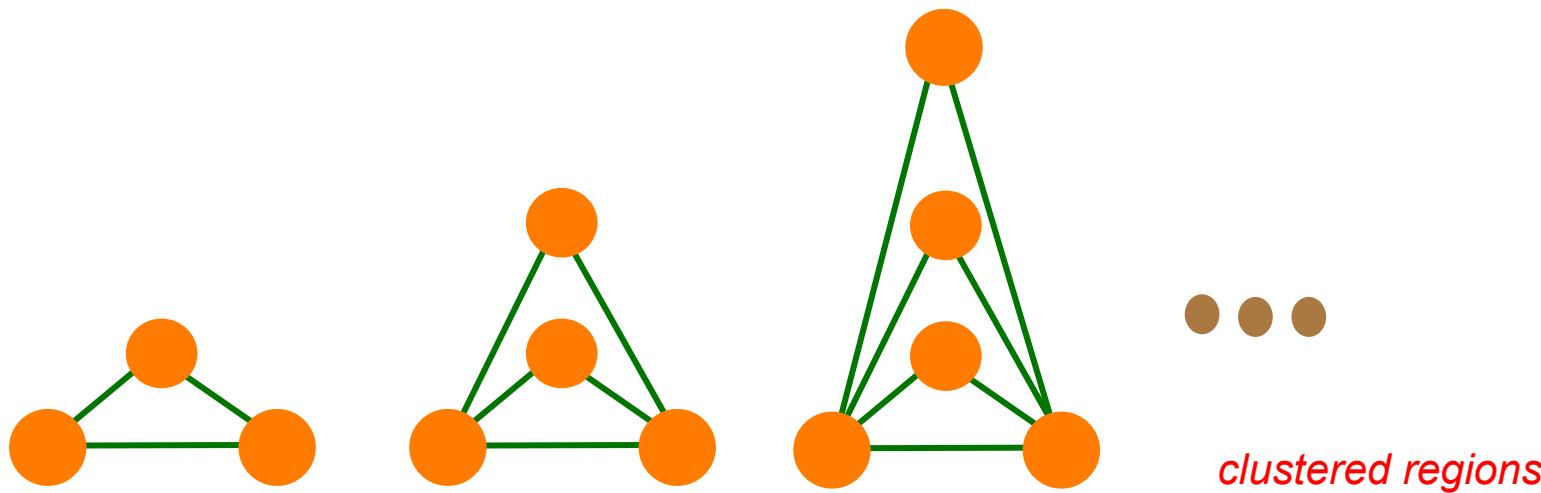
*Distance/similarity*

# ERGMS – modelling graphs

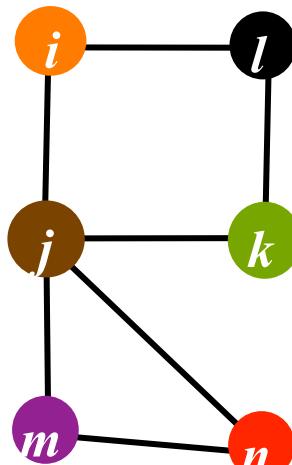
If we believe that (Snijders, et al., 2006)



two edge indicators  $\{i,k\}$  and  $\{l,j\}$  are conditionally  
**dependent** if  $\{i,l\}, \{l,j\} \in E$

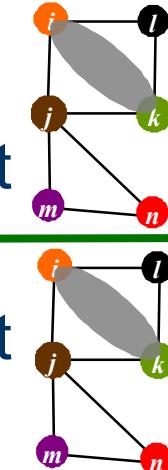


# ERGMS – modelling graphs

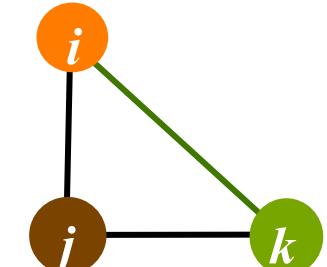
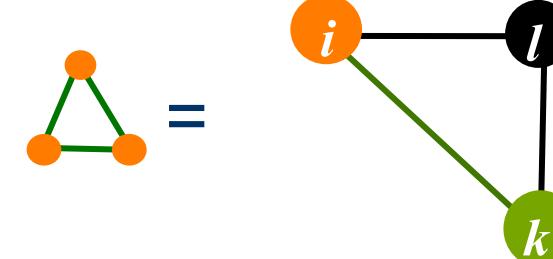
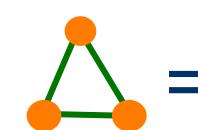


$\Pr \text{ ( } i \text{ --- } k \text{ ) given the rest}$

$\Pr \text{ ( } i \text{ and } k \text{ ) given the rest}$



adding edge, e.g.:  $+ \tau 2 \times$



# ERGMS – modelling graphs

The conditional formulation

$$\log \frac{\Pr \begin{array}{c} i \\[-1ex] \text{---} \\ k \end{array} \text{ given the rest}}{\Pr \begin{array}{c} i \\[-1ex] \text{---} \\ k \end{array} \text{ given the rest}} = \theta_1 \delta_{ik}^1(x) + \theta_2 \delta_{ik}^2(x) + \cdots + \theta_p \delta_{ik}^p(x)$$

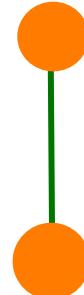
where  $\delta_{ik}^r(x) = z_r(\Delta_{ij}x) - z_r(x)$  Is the **difference in counts of structure type  $k$**

May be "aggregated" for all dyads so that the model for the entire adjacency matrix can be written

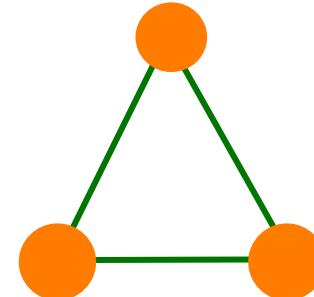
$$\log \Pr(X = x) = \theta_1 z_1(x) + \theta_2 z_2(x) + \cdots + \theta_p z_p(x) + \psi(\theta)$$

## ERGMS – modelling graphs

For a model with edges

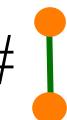
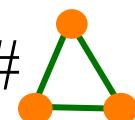


and triangles



The model for the adjacency matrix  $X$  is a weighted sum

$$\log \Pr(X = x) = \sigma_1 L(x) + \tau T(x) + \psi(\theta)$$

where  $L(x) = \#$    $T(x) = \#$  

The parameters  $\sigma_1$  and  $\tau$  weight the relative importance of ties and triangles, respectively

- graphs with many triangles but not too dense are more probable than dense graphs with few triangles

## Padgett's Florentine families (Padgett and Ansell, 1993) network

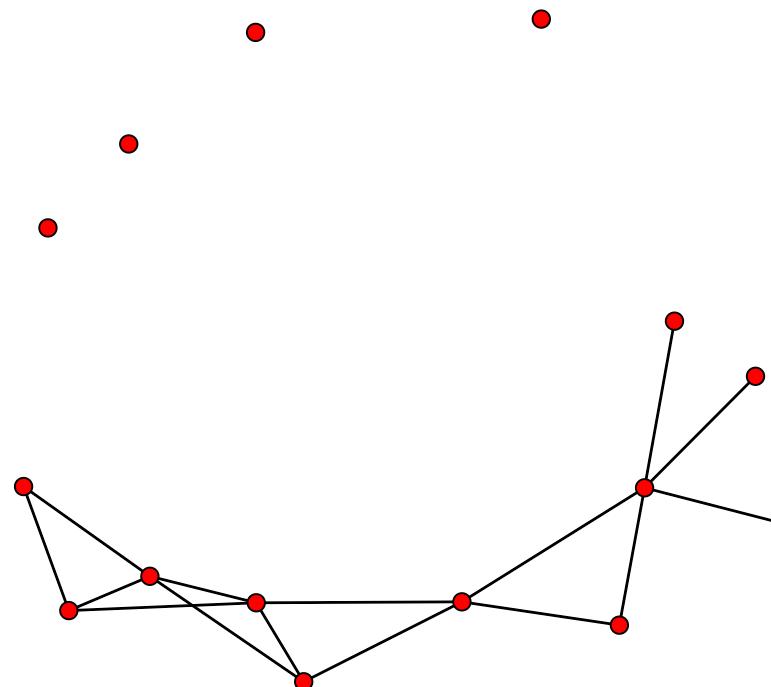
```
BusyNet <- as.matrix(read.table(  
    "PADGB.txt", header=FALSE))
```

```
> BusyNet  
   V1 V2 V3 V4 V5 V6 V7 V8 V9 V10 V11 V12 V13 V14 V15 V16  
[1,] 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0  
[2,] 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0  
[3,] 0 0 0 0 1 1 0 0 1 0 1 0 0 0 0 0 0  
[4,] 0 0 0 0 0 0 1 1 0 0 1 0 0 0 0 0 0  
[5,] 0 0 1 0 0 0 0 1 0 0 1 0 0 0 0 0 0  
[6,] 0 0 1 0 0 0 0 0 1 0 0 0 0 0 0 0 0  
[7,] 0 0 0 1 0 0 0 0 1 0 0 0 0 0 0 0 0  
[8,] 0 0 0 1 1 0 1 0 0 0 1 0 0 0 0 0 0  
[9,] 0 0 1 0 0 1 0 0 0 1 0 0 0 0 1 0 1  
[10,] 0 0 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0  
[11,] 0 0 1 1 1 0 0 1 0 0 0 0 0 0 0 0 0  
[12,] 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0  
[13,] 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0  
[14,] 0 0 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0  
[15,] 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0  
[16,] 0 0 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0  
>
```

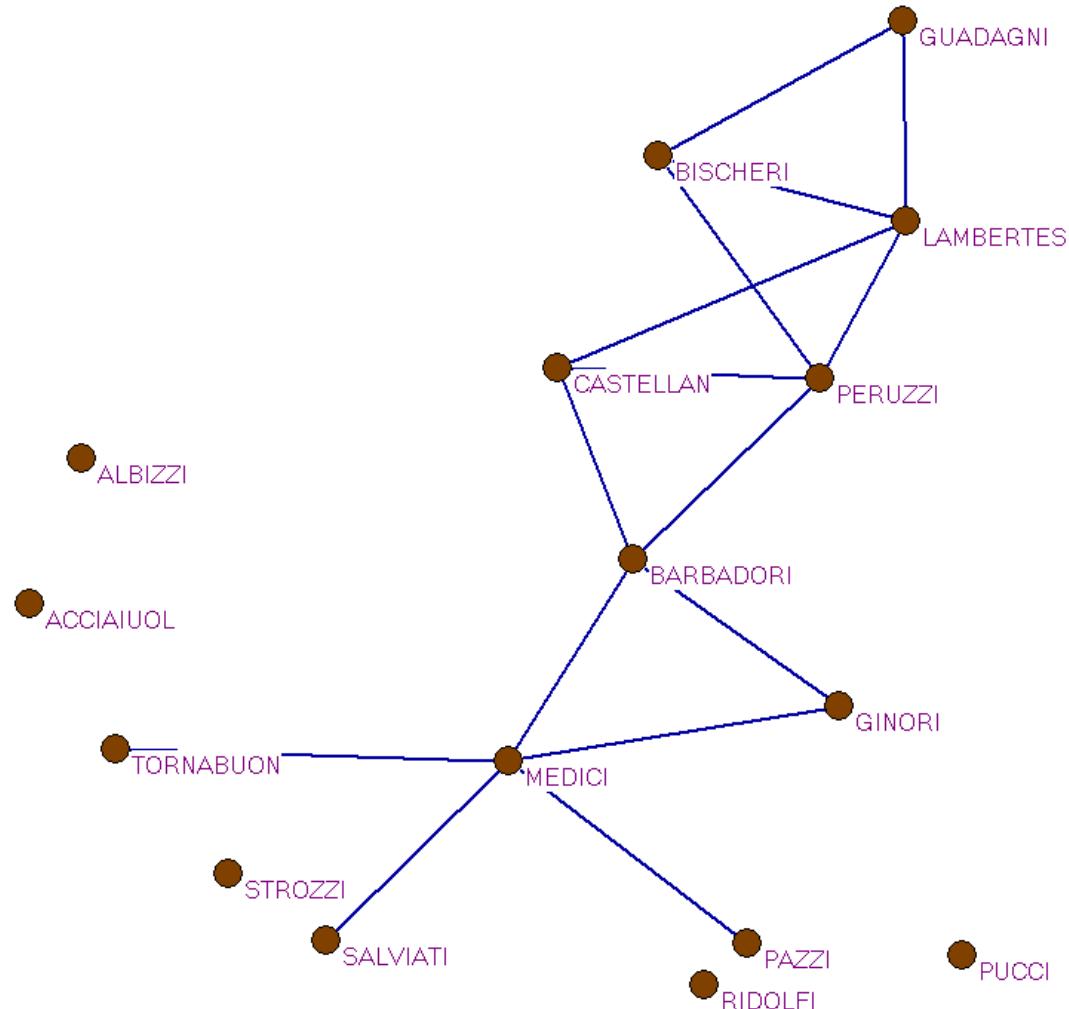
# ERGMS – modelling graphs: example

```
BusyNetNet <- network(BusyNet, directed=FALSE)  
plot(BusyNetNet)
```

Requires libraries  
**'sna'**, **'network'**



# ERGMS – modelling graphs: example



Observed

$$L(x) = \sum_{i < j} x_{ij} = 15$$

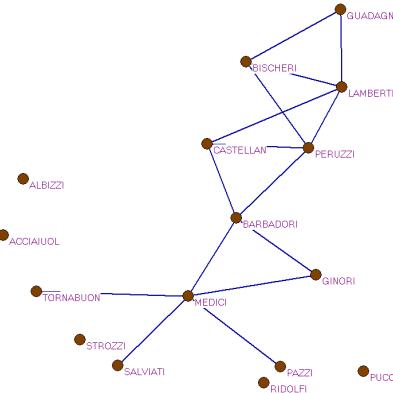
and

$$M = \binom{16}{2} = 120$$

Density:

$$d(x) = \frac{L(x)}{M} = \frac{15}{120} = \frac{1}{8} = 0.125$$

# ERGMS – modelling graphs: example



For a model with only edges



$$\log \Pr(X = x) = \sigma_1 L(x) + \psi(\sigma_1)$$

Equivalent with the model:

For each pair



flip a  $p$ -coin



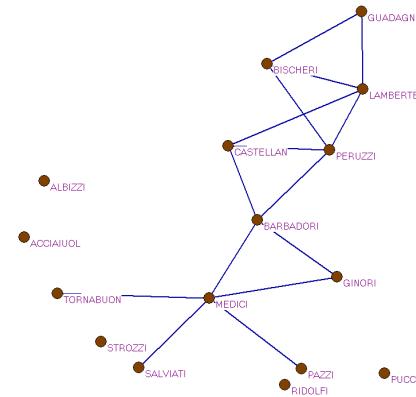
heads

tails



Where  $p$  is the probability coin comes up heads

# ERGMS – modelling graphs: example



The (Maximul likelihood) estimate of  $p$  is the density here:

$$\hat{p}_{MLE} = d(x) = 0.125$$



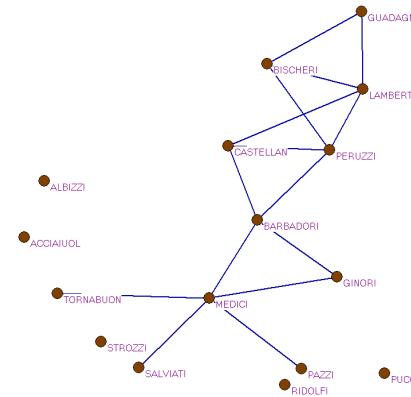
i.e., an estimated one in 8 pairs establish a tie



For an ERGM model with edges  $\log \Pr(X = x) = \sigma_1 L(x) + \psi(\sigma_1)$

and hence 
$$p = \frac{e^{\sigma_1 L(x)}}{1 + e^{\sigma_1 L(x)}}$$

# ERGMS – modelling graphs: example



Solving

$$p = \frac{e^{\sigma_1 L(x)}}{1 + e^{\sigma_1 L(x)}}$$

for  $\sigma_1$ , we have that the density parameter



$$\hat{p}_{MLE} = d(x) = 0.125$$

$$\sigma_1 = -\log(1/p - 1)$$

and for the MLE

$$\hat{\sigma}_{1,MLE} = -\log(1/\hat{p}_{MLE} - 1)$$

$$= -\log(8/1 - 1) = -1.945$$

**Let's check in stanet**

# ERGMS – modelling graphs: example



```
Estim1 <- ergm(BusyNetNet ~ edges)
summary(Estim1)
```

Summary of model fit

Formula: BusyNetNet ~ edges

Newton-Raphson iterations: 5

Maximum Likelihood Results:

	Estimate	Std. Error	MCMC s.e.	p-value
edges	-1.946	0.276	NA	<1e-04 ***

---

Signif. codes: 0 ‘\*\*\*’ 0.001 ‘\*\*’ 0.01 ‘\*’ 0.05 ‘.’ 0.1 ‘ ’ 1

For this model, the pseudolikelihood is the same as the likelihood.

$$\hat{p}_{MLE} = d(x) = 0.125$$

$$\hat{\sigma}_{1,MLE} = -1.945$$

approx. standard error of MLE of  $\sigma_1$ ,

$$\hat{\sigma}_{1,MLE}$$

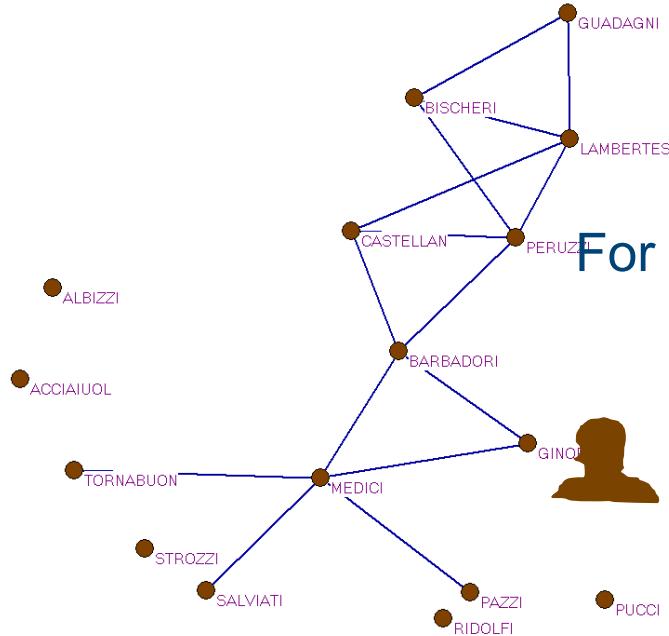
Parameter corresponding to  $L(x) = \#$



**Success:**

$$\hat{\sigma}_{1,MLE} = -\log(1/\hat{p}_{MLE} - 1)$$

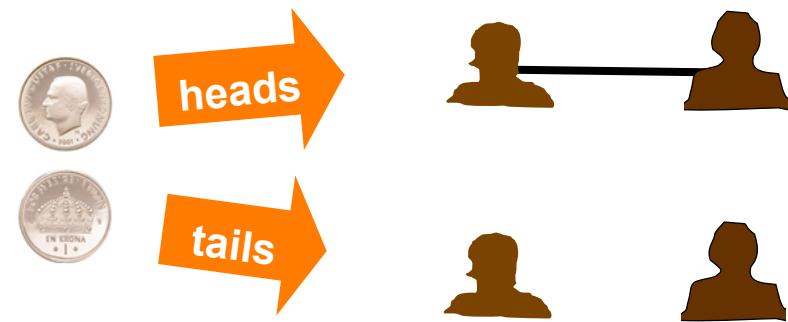
# ERGMS – modelling graphs: example



For each pair

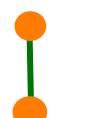
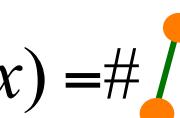
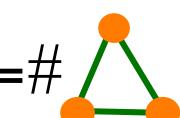
Do we **believe** in the model:

flip a  $p$ -coin



Let's fit a model that takes Markov dependencies into account

$$\log \Pr(X = x) = \sigma_1 L(x) + \sigma_2 S_2(x) + \sigma_3 S_3(x) + \tau T(x) + \psi(\theta)$$

where  $L(x) = \#$    $S_2(x) = \#$    $S_3(x) = \#$    $T(x) = \#$  

**statnet**

# ERGMS – modelling graphs: example

$$L(x) = \# \text{ (isolated node)} \quad S_2(x) = \# \text{ (two nodes connected by one edge)} \quad S_3(x) = \# \text{ (three nodes in a triangle)} \quad T(x) = \# \text{ (triangle)}$$

```
Estim2 <- ergm(BusyNetNet ~ kstar(1:3) + triangles)
summary(Estim2)
```

```
-----
Summary of model fit
-----
```

```
Formula: BusyNetNet ~ kstar(1:3) + triangles
```

```
Newton-Raphson iterations: 42
MCMC sample of size 10000
```

Monte Carlo MLE Results:

	Estimate	Std. Error	MCMC s.e.	p-value
kstar1	-1.6130	0.6699	0.462	0.0176 *
kstar2	0.7492	0.6407	0.455	0.2446
kstar3	-0.5408	0.3574	0.225	0.1330
triangle	1.4837	0.4592	0.138	0.0016 **

---

```
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

approx. standard error of MLE of  $\tau$ ,

$\hat{\tau}_{MLE}$

# Part 2

# Estimation

# Likelihood equations for exponential fam

"Aggregated" to a joint model for **entire adjacency matrix**  $X$

$$\log \Pr(X = x) = \theta_1 z_1(x) + \theta_2 z_2(x) + \cdots + \theta_p z_p(x) + \psi(\theta)$$

Sum over **all**  $2^{n(n-1)/2}$  graphs

The MLE solves the equation (cf. Lehmann, 1983):

$$E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$$

**Solving**  $E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$

- Using the cumulant generating function (Corander, Dahmström, and Dahmström, 1998)
- Stochastic approximation (**Snijders, 2002**, based on **Robbins-Monro, 1951**)
- Importance sampling (**Handcock, 2003**; Hunter and Handcock, 2006, based on **Geyer-Thompson 1992**)

# Robbins-Monro algorithm

**Solving**  $E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$

**Snijders, 2002**, algorithm

- Initialisation phase
- Main estimation
- convergence check and cal. of standard errors

**MAIN:**

$$\theta^{(m+1)} = \theta^{(m)} - a_r D_0^{-1} \{z(x_{\theta^{(m)}}) - z(x_{obs})\}$$

Draw using MCMC

# Robbins-Monro algorithm

## Phase 1, Initialisation phase

Find good values of the initial parameter state

$$\theta^{(0)}$$

And the scaling matrix

$$D_0$$

(use the score-based method, Schweinberger & Snijders, 2006)

## Phase 2, Main estimation phase

Iteratively update  $\theta$

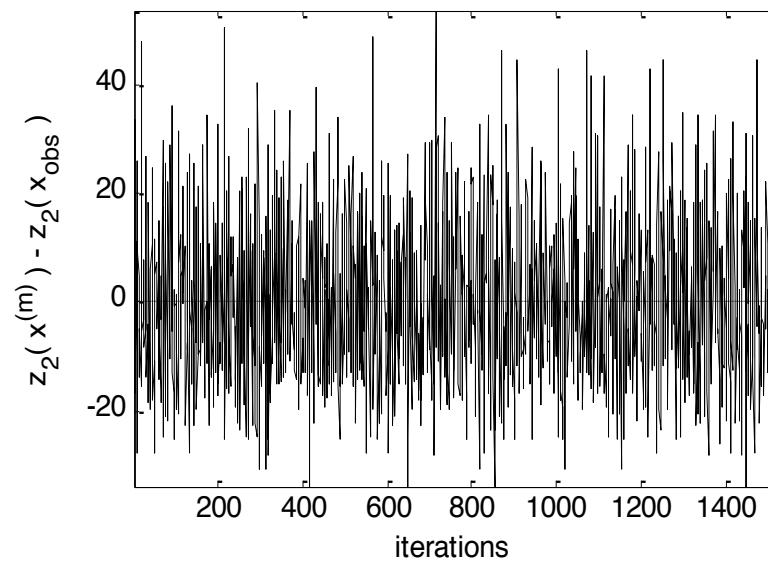
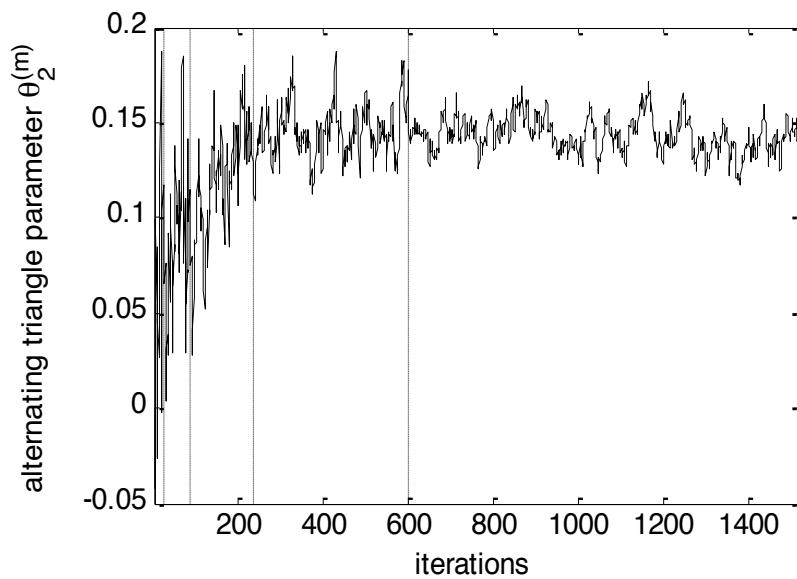
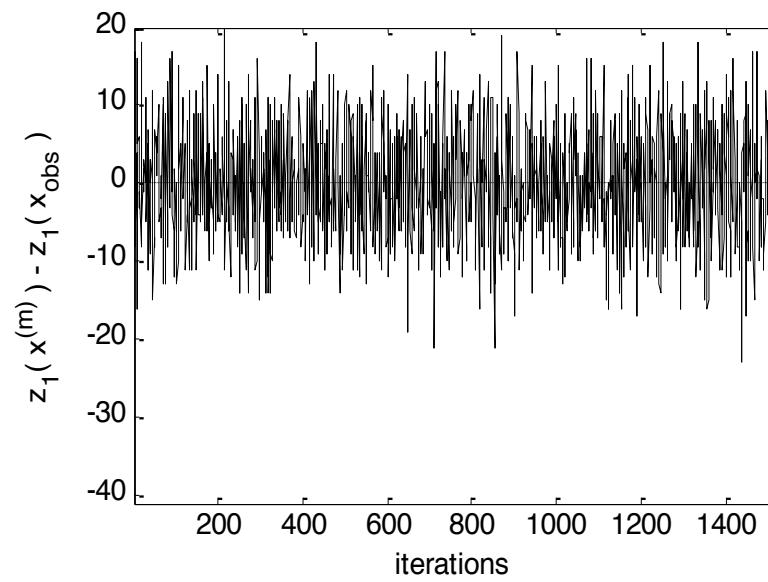
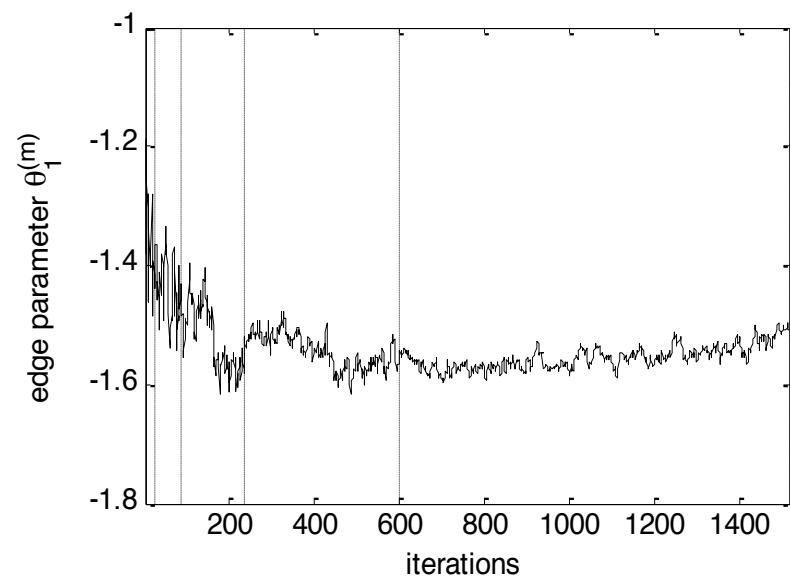
$$\theta^{(m+1)} = \theta^{(m)} - a_r D_0^{-1} \{ z(x_{\theta^{(m)}}^{(m)}) - z(x_{\text{obs}}) \}$$

by drawing one realisation

$$x_{\theta^{(m)}}^{(m)}$$

from the model defined by the current  $\theta^{(m)}$

Repeated in **sub-phases** with fixed  $a_r$



## Robbins-Monro algorithm

### Phase 2, Main estimation phase

Relies on us being able to **draw** one realisation

$x$

from the **ERGM** defined by the current  $\theta$

We can **NOT** do this directly

We have to simulate  $x$

More specifically use **Markov chain Monte Carlo**

## What do we need to know about MCMC?

Method:

Generate a sequence of graphs

$$x^{(0)}, x^{(1)}, x^{(2)}, x^{(3)}, \dots$$

for arbitrary  $x^{(0)}$ , using an updating rule... so that

$$p(x^{(N)}) = \frac{e^{\theta^T z(x^{(N)})}}{\sum_y e^{\theta^T z(y)}} \quad \text{as} \quad N \rightarrow \infty$$

## Robbins-Monro algorithm

### What do we need to know about MCMC?

So if we generate an **infinite** number of graphs in the “**right**” way we have the **ONE** draw we need to update  $\theta$  once?

Typically we can’t wait an infinite amount of time so we settle for

$N \rightarrow$  very large

In Pnet very large is

$$\gamma \text{density}(x_{\text{obs}})[1 - \text{density}(x_{\text{obs}})]n^2$$

multiplication factor

### Phase 3, Convergence check and calculating standard errors

At the end of phase 2 we always get a value

$$\hat{\theta}$$

But is it the MLE?

Does it satisfy

$$E_{\hat{\theta}}\{z(X)\} = z(x_{obs}) \quad ?$$

### Phase 3, Convergence check and calculating standard errors

Phase 3 simulates a large number of graphs  
And checks if

$$E_{\hat{\theta}}\{z(X)\} \approx \overline{E_{\hat{\theta}}\{z(X)\}} \approx z(x_{obs})$$

A minor discrepancy - due to numerical inaccuracy  
- is acceptable

**Convergence statistics:**

$$-.1 < \left| \frac{\overline{E_{\hat{\theta}}\{z(X)\}} - z(x_{obs})}{SD_{\hat{\theta}}\{z(X)\}} \right| < .1$$

**Solving**  $E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$

**Handcock, 2003, approximate Fisher scoring**

**MAIN:**

$$\theta^{(g)} = \theta^{(g-1)} - I(\theta^{(g-1)})^{-1} \left\{ \sum_{m=1}^M w^{(m)} z(x^{(m)}) - z(x_{obs}) \right\}$$

Approximated using importance sample from MCMC

# Bayes: dealing with likelihood

The **normalising constant of the posterior** not essential for Bayesian inference, all we need is:

$$\pi(\theta | x) = \frac{\ell(\theta; x)\pi(\theta)}{\int \ell(\theta; x)\pi(\theta) d\theta} \propto \ell(\theta; x)\pi(\theta)$$

... but

$$\ell(\theta; x) = \frac{\exp\left\{ \sum_{k=1}^p \theta_k z_k(x) \right\}}{\sum \exp\left\{ \sum_{k=1}^p \theta_k z_k(y) \right\}}$$

**Sum over all  $2^{n(n-1)/2}$  graphs**

# Bayes: MCMC?

Consequently,  
in e.g. Metropolis-Hastings, acceptance probability of move to  $\theta$

$$\min\left\{1, \frac{\pi(\theta^* | x) q_{prop}(\theta | \theta^*)}{\pi(\theta | x) q_{prop}(\theta^* | \theta)}\right\} = \min\left\{\frac{\ell(\theta^*; x)\pi(\theta^*)}{\ell(\theta; x)\pi(\theta)} \frac{q_{prop}(\theta | \theta^*)}{q_{prop}(\theta^* | \theta)}\right\}$$

... which contains

$$\frac{\sum_y \exp\{\sum_{k=1}^p \theta_k z_k(y)\}}{\sum_y \exp\{\sum_{k=1}^p \theta_k^* z_k(y)\}}$$

# Bayes: Linked Importance Sampler Auxiliary Variable MCMC

LISA (Koskinen, 2008; Koskinen, Robins & Pattison, 2010): Based on Møller et al. (2006), we define an **auxiliary variable**  $\omega$

$$\omega \in \prod_{j=1}^m \mathcal{X}^K \times \{1, \dots, K\} \times \{1, \dots, K\}$$

And produce draws from the joint posterior

$$\pi(\omega, \theta | x_{obs}) \propto \frac{\exp\{\sum \theta_k z_k(x_{obs})\}}{\sum \exp\{\sum \theta_k z_k(y)\}} \frac{P_{\tau, \theta}^B(\omega)}{\sum \exp\{\sum \tau_k z_k(y)\}} \pi(\theta)$$

using the proposal distributions

$$\theta^* | \theta^{(t)} \sim N(\theta^{(t)}, \Sigma) \quad \text{and} \quad \omega^* | \theta^* \sim \frac{P_{\theta^*, \tau}^F(\omega^*)}{\sum \exp\{\sum \theta_k^* z_k(y)\}}$$

## Bayes: alternative auxiliary variable

LISA (Koskinen, 2008; Koskinen, Robins & Pattison, 2010): Based on Møller et al. (2006), we define an auxiliary variable  $\omega$

$$\omega \in \prod_{j=1}^m \mathcal{X}^K \times \{1, \dots, K\} \times \{1, \dots, K\}$$

Many linked chains:

- Computation time
- storage (memory and time issues)

Improvement: use **exchange algorithm** (Murray et al. 2006)

$$\theta^* | \theta^{(t)} \sim N(\theta^{(t)}, \Sigma) \quad \text{and} \quad x^* | \theta^* \sim \text{ERGM}(\theta^*)$$

Accept  $\theta^*$  with log-probability:

$$\min \{0, (\theta - \theta^*)^T (z(x^*) - z(x_{\text{obs}}))\}$$

# Bayes: Implications of using alternative auxiliary variable

- Storing only parameters
- No pre tuning – no need for good initial values
- Standard MCMC properties of sampler
- Less sensitive to near degeneracy in estimation
- Easier than anything else to implement

**QUICK** and **ROBUST**

Improvement: use **exchange algorithm** (Murray et al. 2006)

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Accept  $\theta^*$  with log-probability:

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## Bayes: Implications of using alternative auxiliary variable

**exchange algorithm** (Murray et al. 2006)

auxiliary variables:  $h(\theta^* | \theta)$

and  $p(x^* | \theta^*) \sim \text{ERGM}(\theta^*)$

## Bayes: Implications of using alternative auxiliary variable

exchange algorithm (Murray et al. 2006)

auxiliary variables:  $h(\theta^* | \theta)$

and  $p(x^* | \theta^*) \sim \text{ERGM}(\theta^*)$

To draw from joint posterior  $\propto p(x^* | \theta^*)h(\theta^* | \theta)p(x | \theta)\pi(\theta)p(x | \theta)$

## Bayes: Implications of using alternative auxiliary variable

**exchange algorithm** (Murray et al. 2006)

**auxiliary** variables:  $h(\theta^* | \theta)$

and  $p(x^* | \theta^*) \sim \text{ERGM}(\theta^*)$

To draw from **joint** posterior  $\propto p(x^* | \theta^*)h(\theta^* | \theta)p(x | \theta)\pi(\theta)p(x | \theta)$

**Gibbs-draw:**  $(x^* | \theta^*) \sim p(x^* | \theta^*)h(\theta^* | \theta)$

# Bayes: Implications of using alternative auxiliary variable

**exchange algorithm** (Murray et al. 2006)

**auxiliary** variables:  $h(\theta^* | \theta)$

and  $p(x^* | \theta^*) \sim \text{ERGM}(\theta^*)$

To draw from **joint** posterior  $\propto p(x^* | \theta^*)h(\theta^* | \theta)p(x | \theta)\pi(\theta)p(x | \theta)$

**Gibbs-draw:**  $(x^* | \theta^*) \sim p(x^* | \theta^*)h(\theta^* | \theta)$

then swap  $\theta^*$  and  $\theta$  with probability  $\min\{1, H\}$

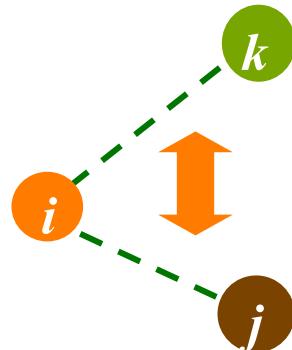
$$H = \frac{p(x_{\text{obs}} | \theta^*)}{p(x_{\text{obs}} | \theta)} \frac{\pi(\theta^*)}{\pi(\theta)} \frac{h(\theta | \theta^*)p(x^* | \theta)}{h(\theta^* | \theta)p(x^* | \theta^*)}$$

## Part 3

### Interpretation of effects

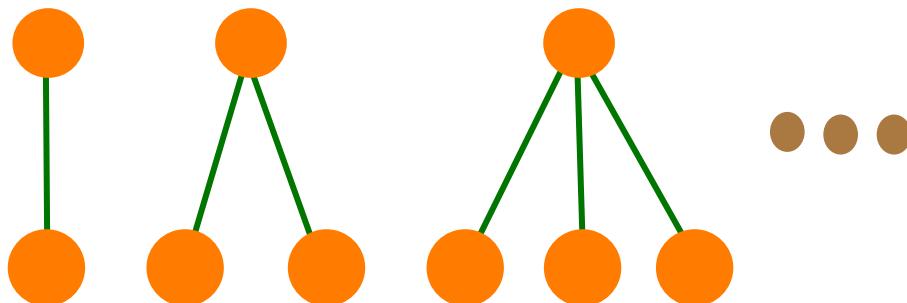
# Problem with Markov models

**Markov dependence** assumption:

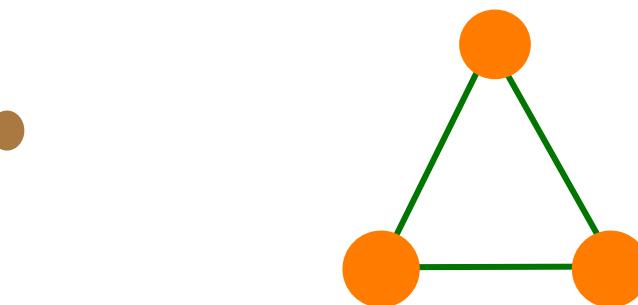


two edge indicators  $\{i,j\}$  and  $\{i',k\}$  are conditionally **dependent** if  $\{i,j\} \cap \{i',k\} \neq \emptyset$

We have shown that the **only** effects are:



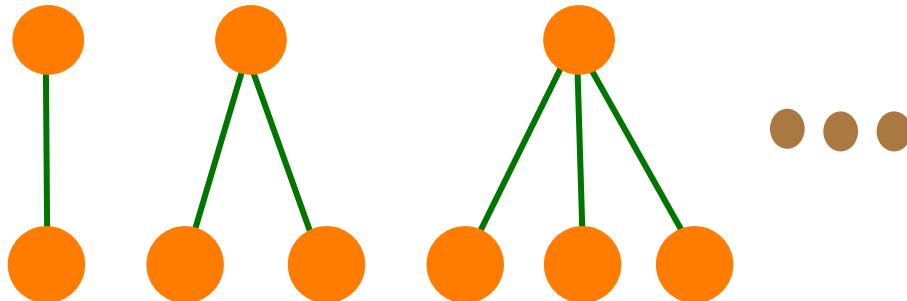
*degree distribution; preferential attachment,  
etc*



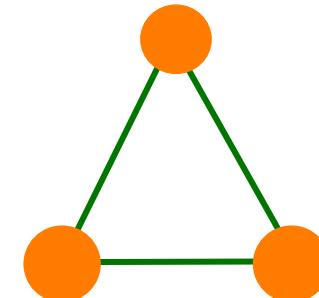
*friends meet through friends; clustering; etc*

# Problem with Markov models

Often for Markov model



*degree distribution; preferential attachment,  
etc*



*friends meet through friends; clustering; etc*

**Matching**

$$E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$$



or



# Problem with Markov models

Matching

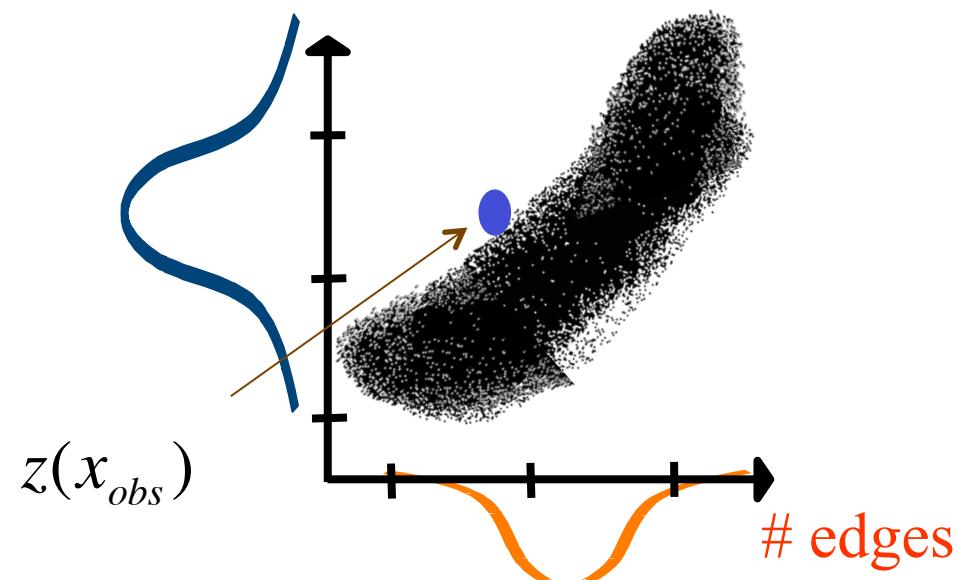
$$E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$$



or

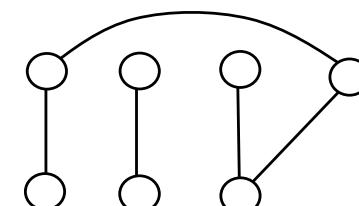


# triangles



Some statistic  $k$

$$z_k(x_{obs}) = 0 \text{ or } z_k(x_{obs}) = z_k^{\max}$$



# Problem with Markov models

**Matching**

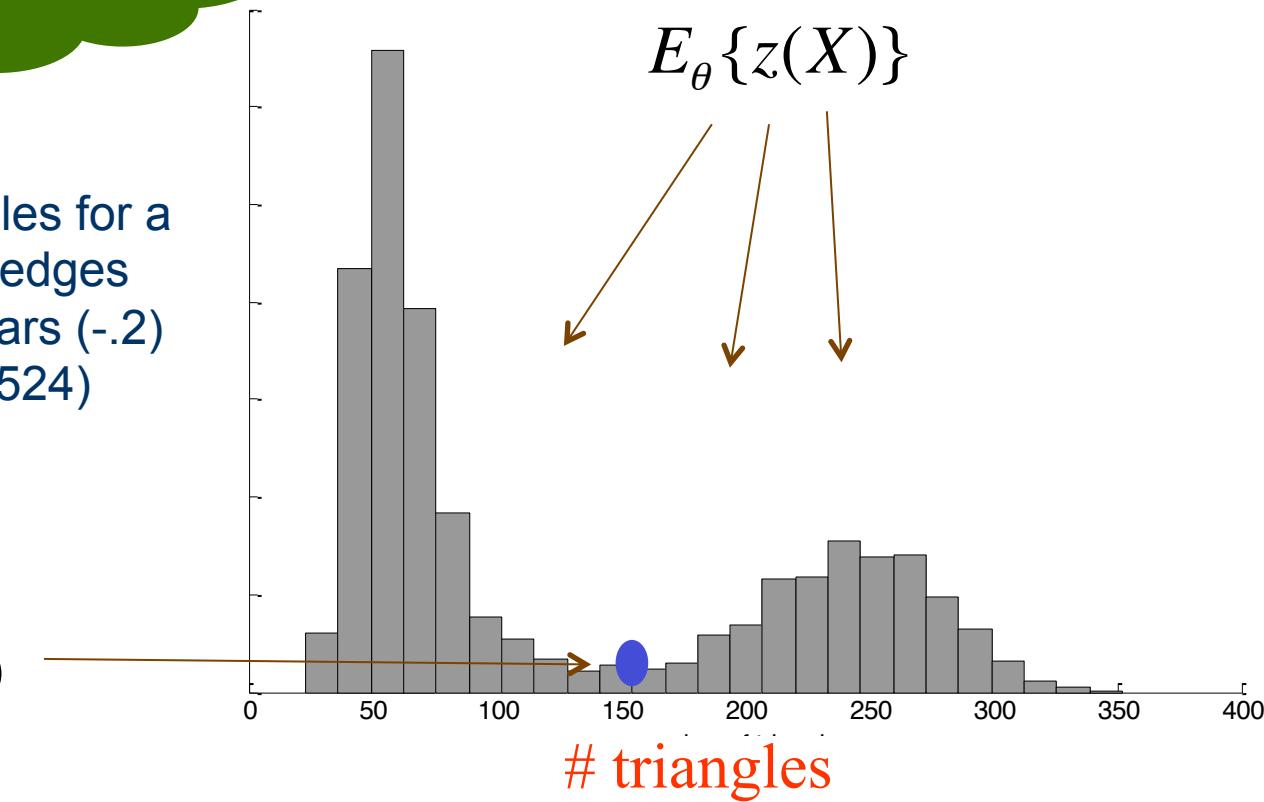
$$E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$$

hard

The number of triangles for a Markov model with edges (-3), 2-stars (.5), 3-stars (-.2) and triangles (1.1524)

$$z(x_{obs})$$

$$E_{\theta} \{z(X)\}$$



# Problem with Markov models

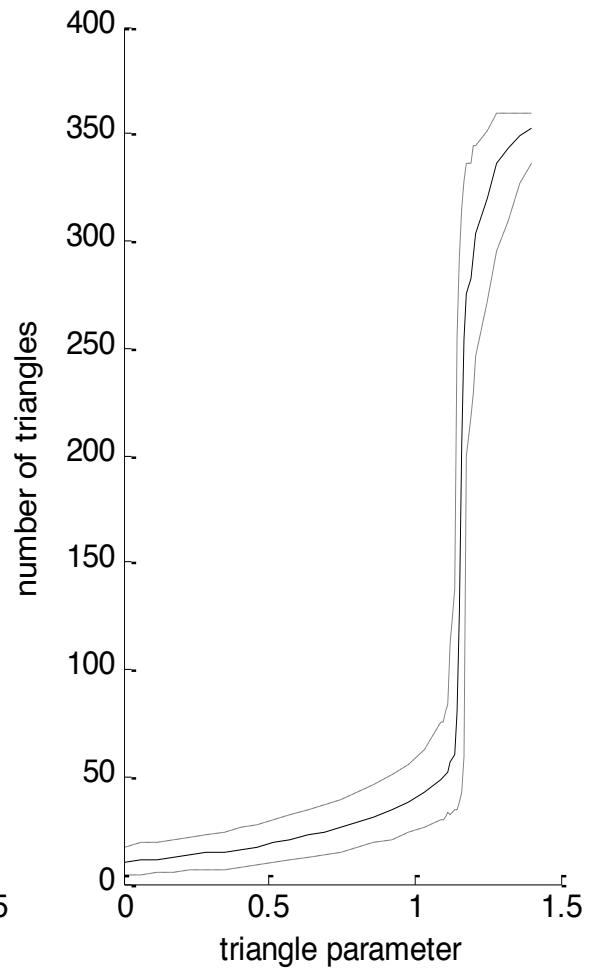
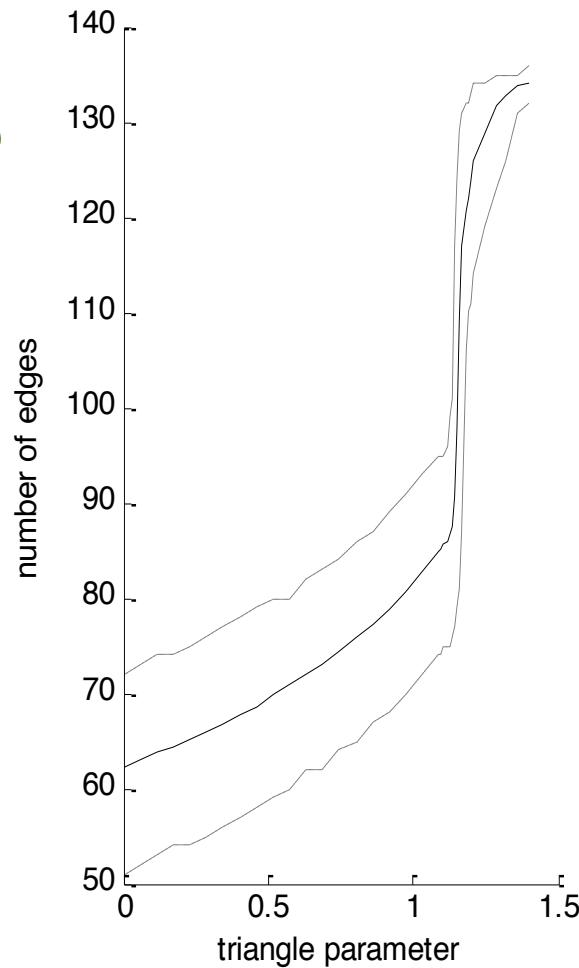
## Matching

$$E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$$

hard

Exp. # triangles and edges for a Markov model with edges, 2-stars, 3-stars and triangles

$$E_{\theta} \{z(X)\}$$



# Problem with Markov models

**Matching**

$$E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$$

Impossible!

If for some statistic  $k$   $z_k(x_{obs}) = 0$

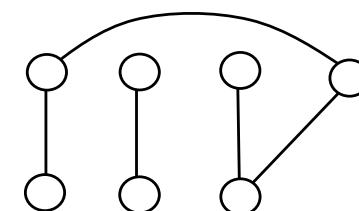
$$E_{\theta} \{z(X)\} = \sum_y z_k(y) p(y | \theta) = z_k(x_{obs})$$

Implies:  $p(y | \theta) = \begin{cases} 1 & \text{if } z_k(y) = z_k(x_{obs}) = 0 \\ 0 & \text{otherwise} \end{cases}$

Similarly for **max**

(or conditional min/max)

e.g. A graph on 7 vertices with 5 edges



# edges      # 2-stars

## Problem with Markov models

Matching

$$E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$$



Impossible!

Generally, let  $C$  be the convex hull of  $\{z(x) : x \in \mathcal{X}\}$

Let  $\text{rint}(C)$  denote the relative interior of  $C$

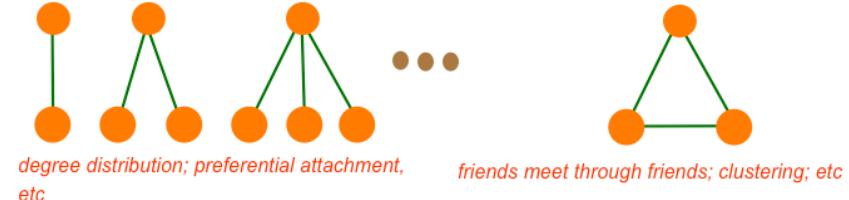
The MLE exists if and only if  $z(x_{obs}) \in \text{rint}(C)$

With constant prior, the posterior exists  
only if  $z(x_{obs}) \in \text{rint}(C)$

See Handcock (2003)

# Problem with Markov models

First **solutions**  
(Snijders et al., 2006)



**Markov:**

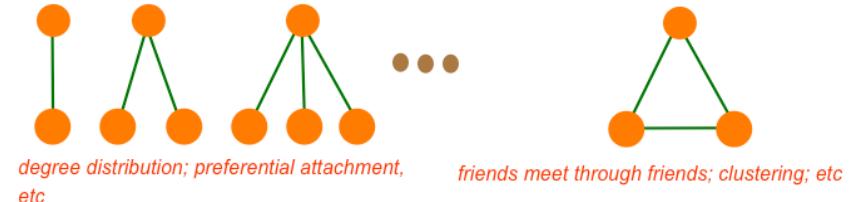
- adding  $k$ -star adds  $(k-1)$  stars
- alternating sign compensates but eventually  $z_k(x)=0$

**Alternating stars:**

- Restriction on star parameters - Alternating sign
- prevents explosion, and
- models degree distribution

# Problem with Markov models

First **solutions**  
(Snijders et al., 2006)



**Markov:**

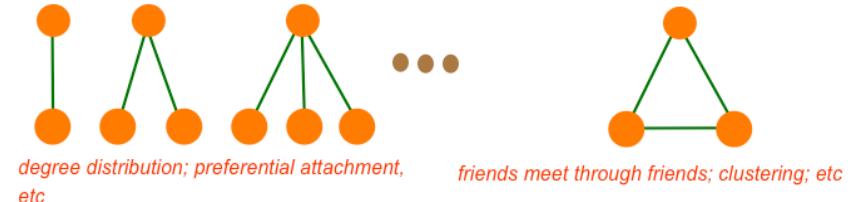
- adding  $k$ -star adds  $(k-1)$  stars
- alternating sign compensates but eventually  $z_k(x)=0$

	Estimate	Std. Error	MCMC s.e.	p-value	
kstar1	-1.6130	0.6699	0.462	0.0176	*
kstar2	0.7492	0.6407	0.455	0.2446	
kstar3	-0.5408	0.3574	0.225	0.1330	
triangle	1.4837	0.4592	0.138	0.0016	**

$$L(x) = \# \text{ } \begin{array}{c} \text{---} \\ | \\ \bullet \end{array} \quad S_2(x) = \# \text{ } \begin{array}{c} \text{---} \\ | \\ \bullet \text{---} \bullet \end{array} \quad S_3(x) = \# \text{ } \begin{array}{c} \text{---} \\ | \\ \bullet \text{---} \bullet \text{---} \bullet \end{array} \quad T(x) = \# \text{ } \begin{array}{c} \text{---} \\ | \\ \bullet \text{---} \bullet \text{---} \bullet \end{array}$$

# Problem with Markov models

First **solutions**  
(Snijders et al., 2006)



Include all stars but restrict parameter:

$$\sigma_3 = -\sigma_2 / \lambda \quad \sigma_4 = -\sigma_3 / \lambda \quad \sigma_5 = -\sigma_4 / \lambda \quad \dots$$

new

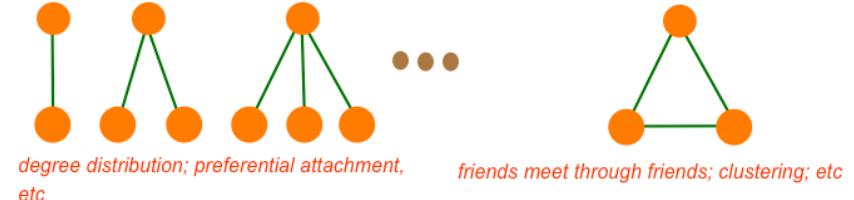
$$\sigma_2 S_2(x) + \sigma_3 S_3(x) + \dots + \sigma_{n-1} S_{n-1}(x) = \sigma_{AKS} AKS(x; \lambda)$$

Alternating stars:

- Restriction on star parameters - Alternating sign
- prevents explosion, and
- models degree distribution

# Problem with Markov models

First **solutions**  
(Snijders et al., 2006)



Include all stars but restrict parameters:

$$\sigma_3 = -\sigma_2 / \lambda \quad \sigma_4 = -\sigma_3 / \lambda \quad \sigma_5 = -\sigma_4 / \lambda \quad \dots$$

new

$$\sigma_2 S_2(x) + \sigma_3 S_3(x) + \dots + \sigma_{n-1} S_{n-1}(x) = \sigma_{AKS} AKS(x; \lambda)$$

Expressed in terms  
of degree distribution

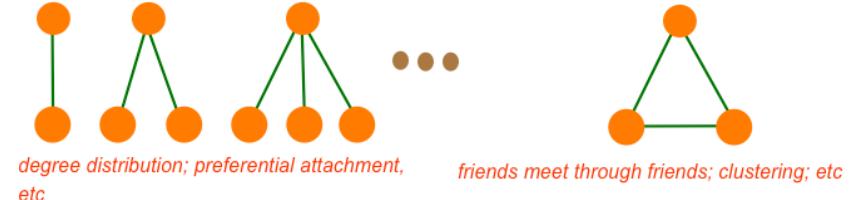
$$= \left( \frac{1}{1 - e^{-\alpha}} \right)^2 \sum_{j=0}^{n-1} d_j(x) e^{-\alpha j} + \frac{2L(x)}{1 - e^{-\alpha}} - \frac{n}{(1 - e^{-\alpha})^2}$$

$$d_j(x) = \#\{i : x_{i+} = j\}$$

$$\lambda = e^\alpha / (e^\alpha - 1)$$

# Problem with Markov models

First **solutions**  
(Snijders et al., 2006)



Include all stars but restrict parameters:

$$\sigma_3 = -\sigma_2 / \lambda \quad \sigma_4 = -\sigma_3 / \lambda \quad \sigma_5 = -\sigma_4 / \lambda \quad \dots$$

$$\sigma_2 S_2(x) + \sigma_3 S_3(x) + \dots + \sigma_{n-1} S_{n-1}(x) = \sigma_{AKS} AKS(x; \lambda)$$

Interpretation:

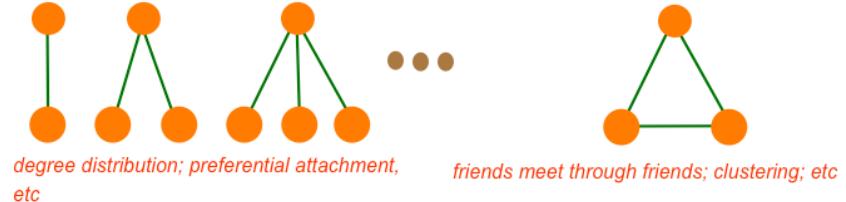
**Positive parameter** ( $\lambda \geq 1$ ) – graphs with some high degree nodes and larger degree variance more likely than graphs with more homogenous degree distribution

**Negative parameter** ( $\lambda \leq 1$ ) – the converse...

# Problem with Markov models

First **solutions**  
(Snijders et al., 2006)  
**Markov:**

- ❖ triangles evenly spread out
- ❖ but one edge can add many triangles...



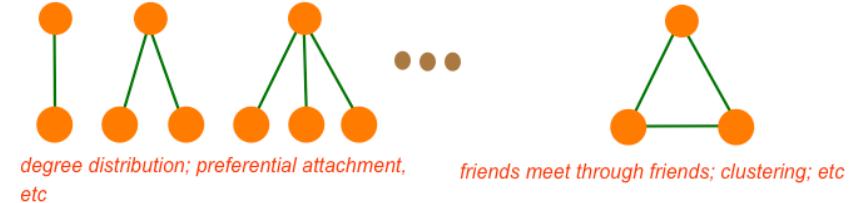
**Alternating triangles:**

- Restrictions on different order triangles – alternating sign
- Prevents explosion, and
- Models multiply clustered regions
- **Social circuit dependence assumption**

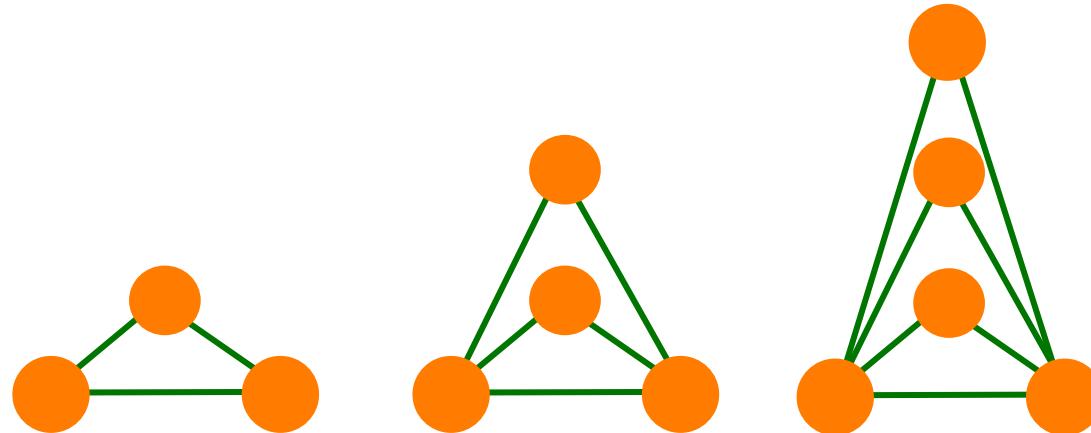
# Problem with Markov models

First solutions  
(Snijders et al., 2006)  
**Markov:**

- ❖ triangles evenly spread out
- ❖ but one edge can add many triangles...



$k$ -triangles:



1-triangles:    2-triangles:    3-triangles:

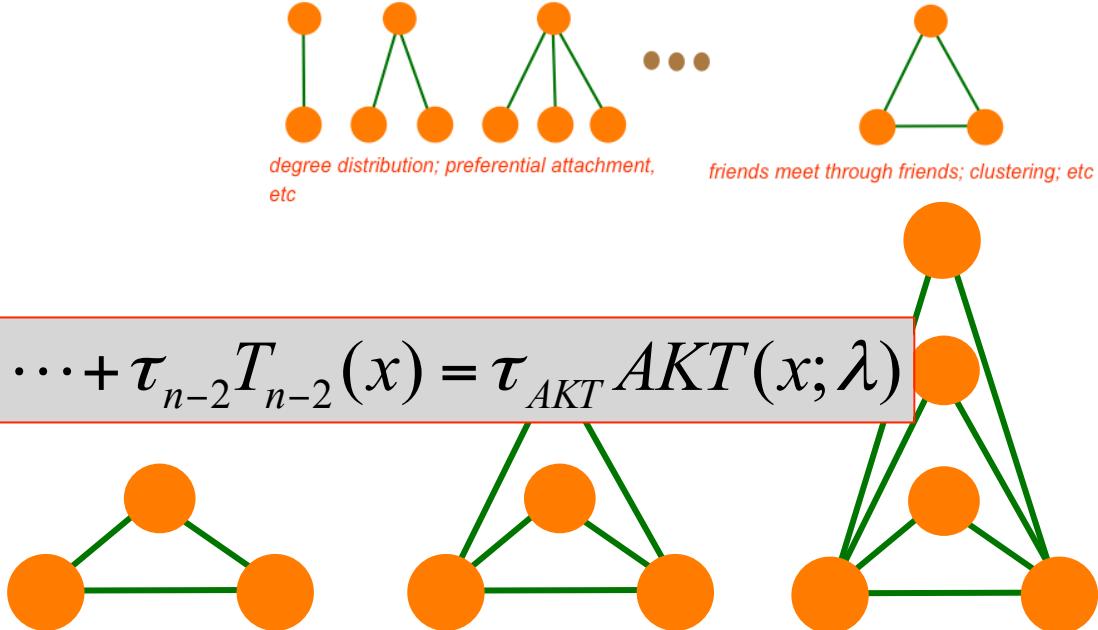
# Problem with Markov models

## First solutions

Weigh the  $k$ -triangles:

$$\tau_2 T_2(x) + \tau_3 T_3(x) + \cdots + \tau_{n-2} T_{n-2}(x) = \tau_{AKT} AKT(x; \lambda)$$

Where:  $\tau_k = -\tau_{k-1} / \lambda$



## Alternating triangles:

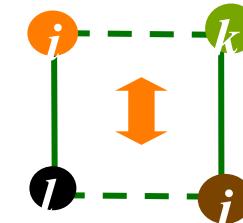
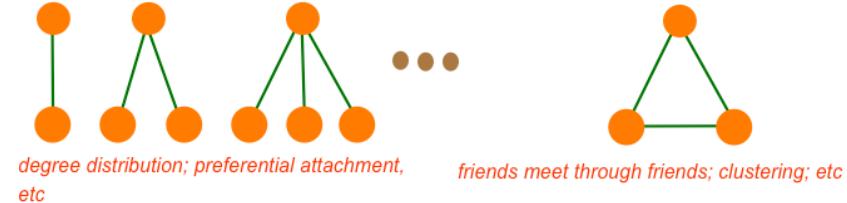
- Restrictions on different order triangles – alternating sign
- Prevents explosion, and
- Models multiply clustered regions

# Problem with Markov models

## First solutions

Underlying assumption:

**Social circuit dependence assumption**



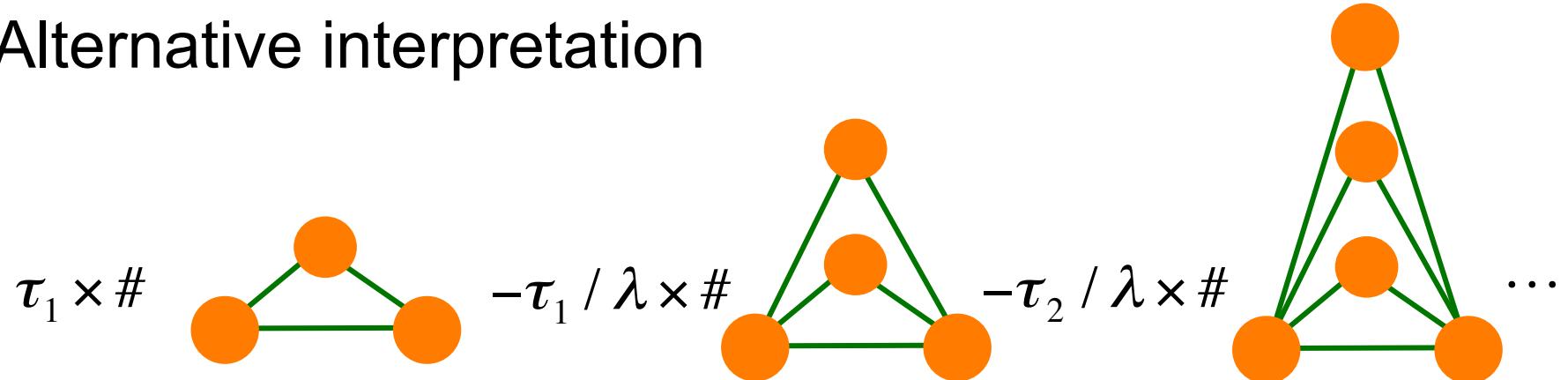
two edge indicators  $\{i,k\}$  and  $\{l,j\}$  are conditionally **dependent** if  $\{i,l\}, \{l,j\} \in E$

**Alternating triangles:**

- Restrictions on different order triangles – alternating sign
- Prevents explosion, and
- Models multiply clustered regions

# Problem with Markov models

Alternative interpretation

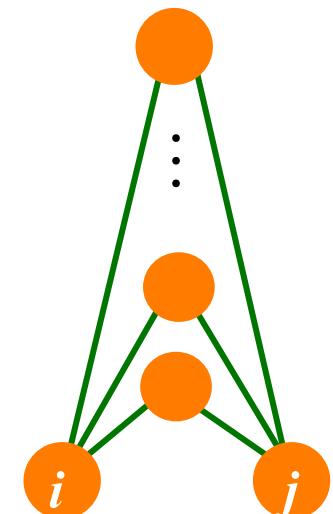


We may (geometrically) weight together :

$$z_T(x; \lambda) = \frac{e^\alpha}{e^\alpha - 1} \left\{ \sum_{i < j} x_{ij} - \sum_{i < j} x_{ij} \frac{1}{e^{\alpha S_{2ij}(x)}} \right\}$$

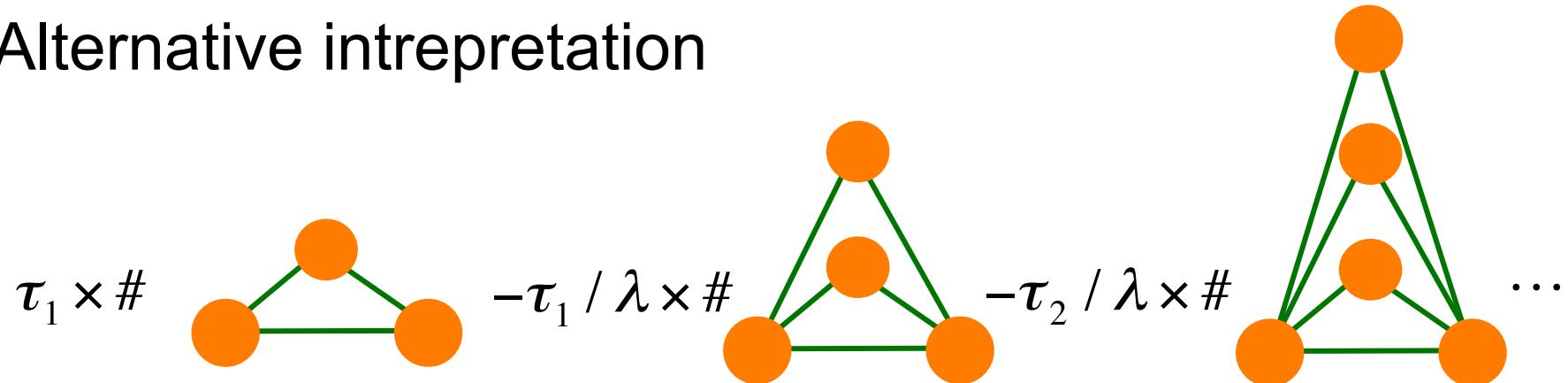
$$\lambda = e^\alpha / (e^\alpha - 1)$$

$$S_{2ij} = \#\{k : i \rightarrow k, j \rightarrow k\}$$



# Problem with Markov models

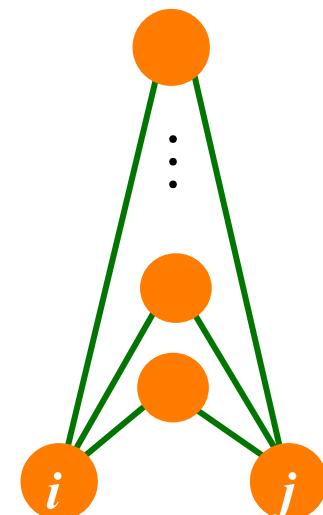
Alternative interpretation



We may also define the  
**Edgewise Shared Partner Statistic:**

$$ESP_k = \#\{(i, j) : i \sim j, S_{2ij} = k\}$$

... and we can weigh together the **ESP** statistics using **Geometrically decreasing weights: GWESP**



## Part 3b

Curved exponential family distributions for graphs

In an ERGM, **alternating** statistics

**alternating stars**

$$z_S(x; \alpha) = \left( \frac{1}{1 - e^{-\alpha}} \right)^2 \sum_{j=0}^{n-1} d_j(x) e^{-\alpha j} + \frac{2L(x)}{1 - e^{-\alpha}} - \frac{n}{(1 - e^{-\alpha})^2}$$

**alternating triangles**

$$z_T(x; \alpha) = \frac{e^\alpha}{e^\alpha - 1} \left\{ \sum_{i < j} x_{ij} - \sum_{i < j} x_{ij} \frac{1}{e^{\alpha S_{2ij}(x)}} \right\}$$

... are “dampened” by a constant  $\alpha$

why not estimate  $\alpha$ ?

## Curved ERGM

If we treat  $\alpha$  as free parameters to be estimated

$$p(x) = \exp\{\theta^T z(x; \alpha) - \psi(\theta, \alpha)\}$$

If we treat  $\alpha$  as free parameters to be estimated

$$p(x) = \exp\{\theta^T z(x; \alpha) - \psi(\theta, \alpha)\}$$

We have **more statistics** than **parameters**  
... it is no longer an exponential family distribution

If we treat  $\alpha$  as free parameters to be estimated

$$p(x) = \exp\{\theta^T z(x; \alpha) - \psi(\theta, \alpha)\}$$

We have **more statistics** than **parameters**  
... it is no longer an exponential family distribution

For example, we no longer have the identity

$$E_{\hat{\theta}_{MLE}}\{z(X)\} = z(x_{obs})$$

## Curved ERGM

If we treat  $\alpha$  as free parameters to be estimated

$$p(x) = \exp\{\theta^T z(x; \alpha) - \psi(\theta, \alpha)\}$$

We have **more statistics** than **parameters**  
... it is no longer an exponential family distribution

However, does not matter for **Bayesian** analysis

$$\pi(\theta, \alpha | x) \propto \exp\{\theta^T z(x; \alpha) - \psi(\theta, \alpha)\} \pi(\theta, \alpha)$$

If we treat  $\alpha$  as free parameters to be estimated

$$p(x) = \exp\{\theta^T z(x; \alpha) - \psi(\theta, \alpha)\}$$

We have **more statistics** than **parameters**

... it is no longer an exponential family distribution

Formally it is a **Curved** exponential family distribution  
... and a Fisher scoring algorithm (using MCMC) can be applied (Hunter and Handcock, 2006)

## Part 4a

Example Lazega's law firm partners

# Lazega's (2001) Lawyers

Collaboration network among 36 lawyers in a  
New England law firm (Lazega, 2001)

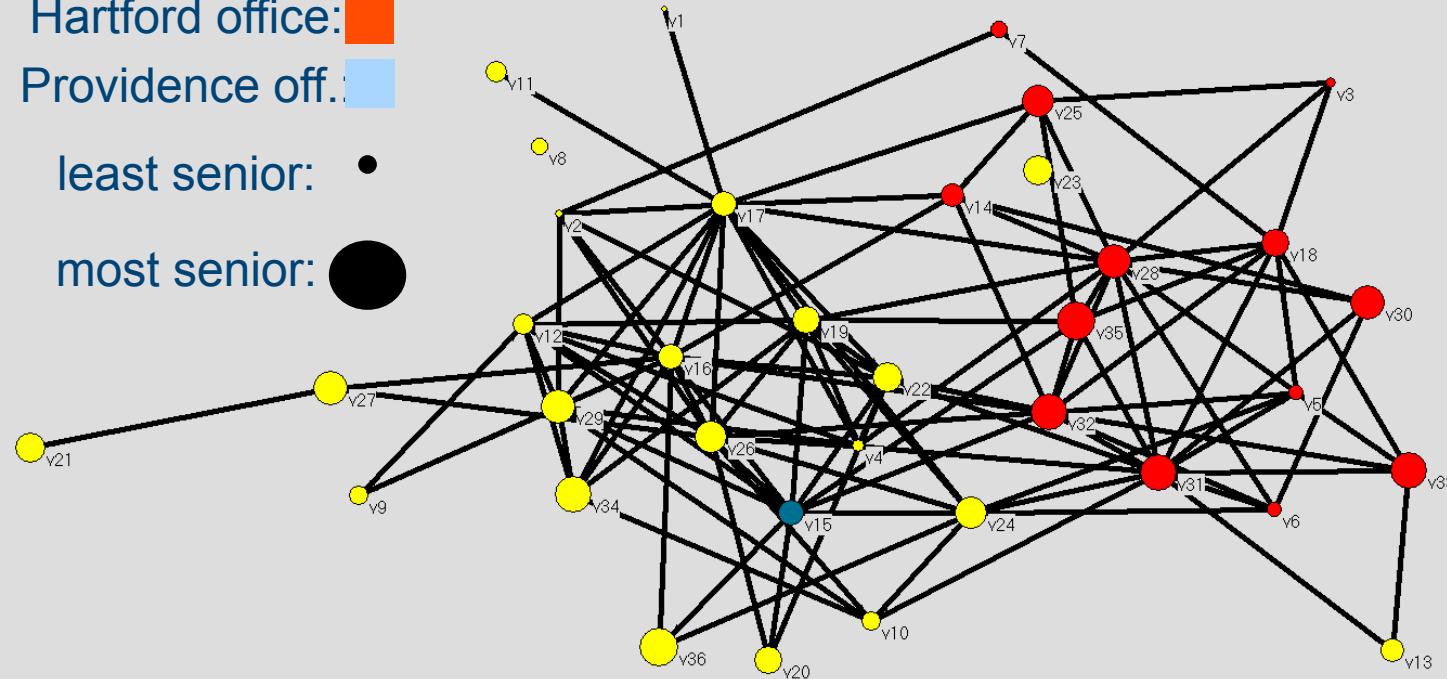
Boston office: ■

Hartford office: □

Providence off.: ▲

least senior: ●

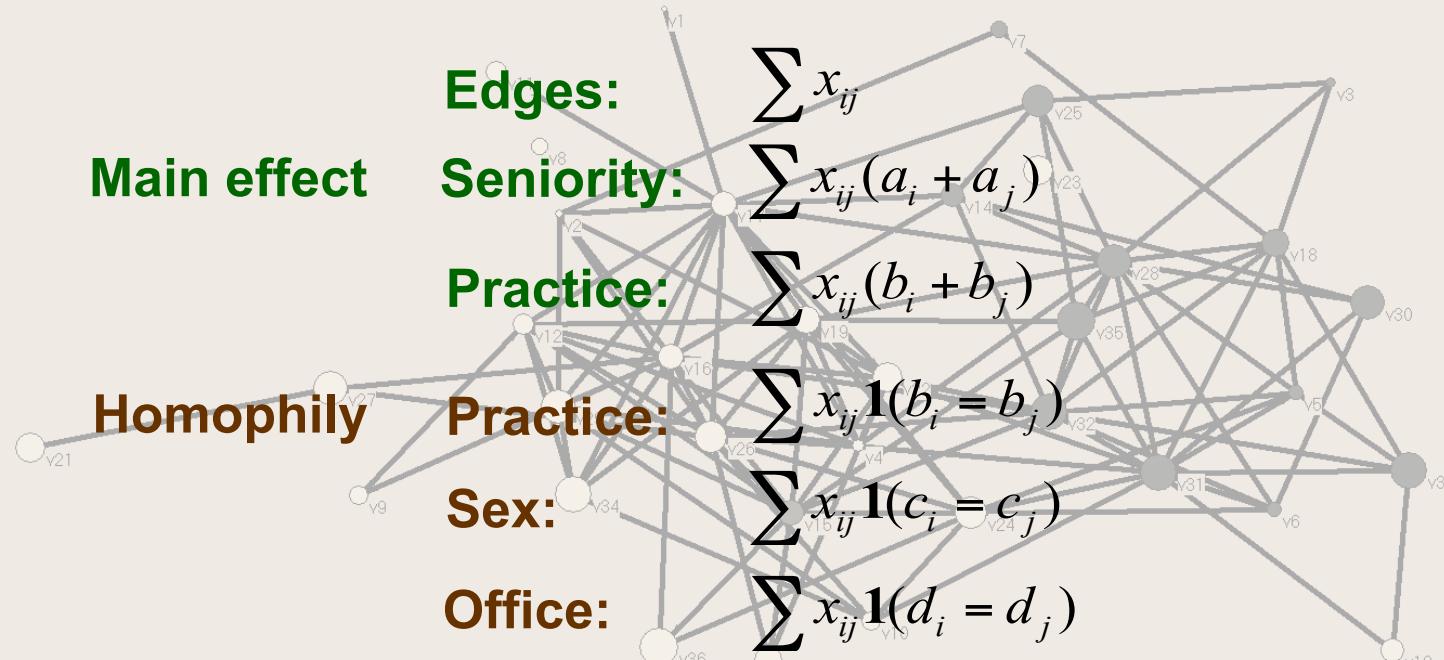
most senior: ●



# Lazega's (2001) Lawyers

Fit a model with "**new specifications**" and **covariates**

$$\log \Pr(X = x) = \theta_1 z_1(x) + \theta_2 z_2(x) + \cdots + \theta_p z_p(x) + \psi(\theta)$$



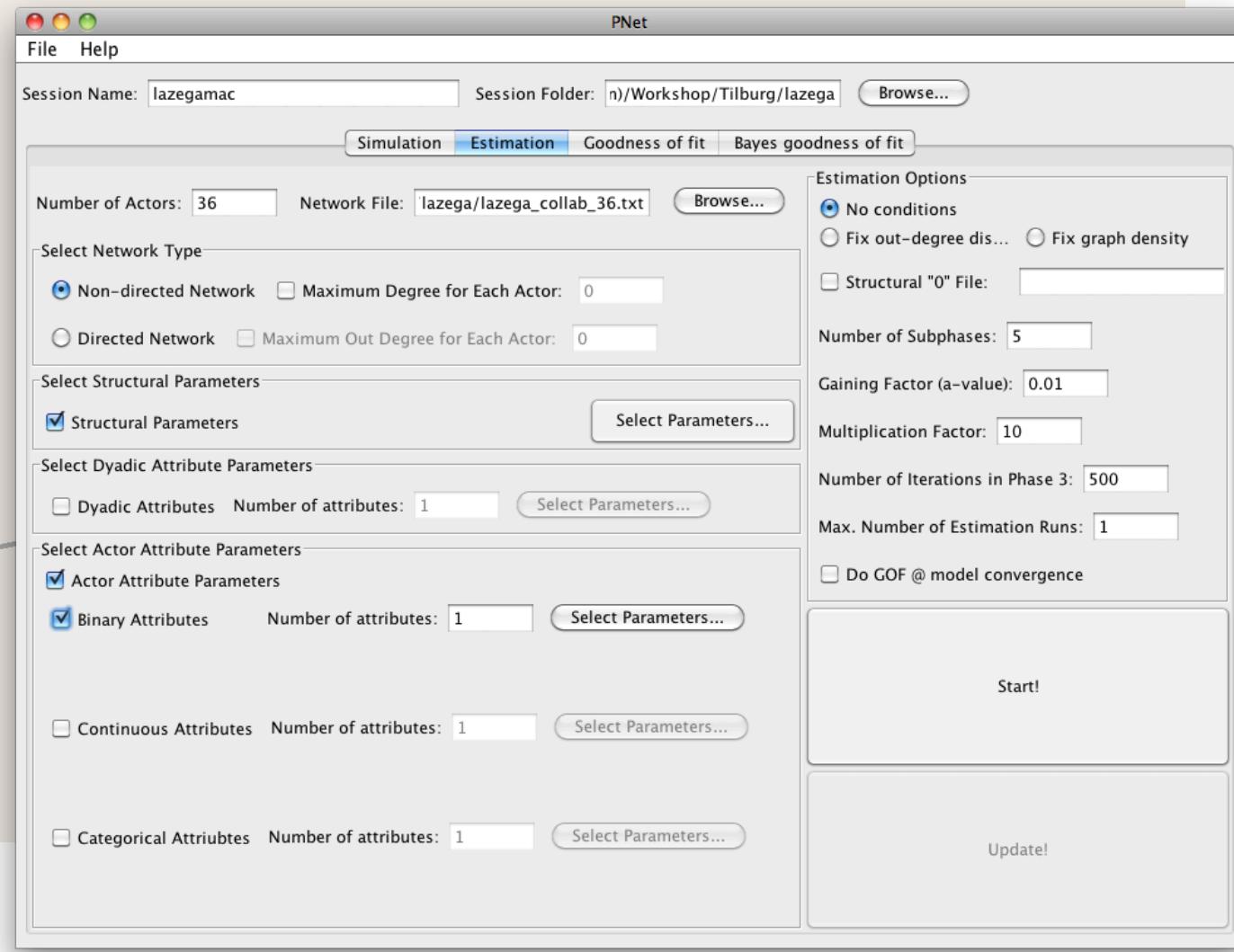
GWESP:  $3t_1(x) - \frac{t_2(x)}{\lambda^1} + \cdots + (-1)^{n-3} \frac{t_{n-2}(x)}{\lambda^{n-3}}$

# Lazega's (2001) Lawyers

Fit a model with "new specifications" and covariates

PNet:

y21



# Lazega's (2001) Lawyers

Fit a model with "new specifications" and covariates

## Main effect

**Edges:**  $\sum x_{ij}$

**Seniority:**  $\sum x_{ij}(a_i + a_j)$

**Practice:**  $\sum x_{ij}(b_i + b_j)$

## Homophily

**Practice:**  $\sum x_{ij} \mathbf{1}(b_i = b_j)$

**Sex:**  $\sum x_{ij} \mathbf{1}(c_i = c_j)$

**Office:**  $\sum x_{ij} \mathbf{1}(d_i = d_j)$

estimation_lazega.txt			
*****			
mean statistics	in phase3:114.362000	179.720562	42.846000
128.654000	129.287641	98.156000	84.984000
Estimation Result for Network SUMMARY (parameter, standard error, t-statistics)			
NOTE: t-statistics = (observation - sample mean)/standard error			
effects	estimates	stderr	t-ratio
edge	-5.862515	0.56404	0.04105 *
AT(2.00)	1.011721	0.17095	0.05003 *
practice_interaction	1.499409	0.40322	0.02371 *
practice_activity	-0.331023	0.21995	0.02142
senior_sum	0.842661	0.23348	0.04943 *
sex_matching	0.702477	0.26389	0.05839 *
off_matching	1.145290	0.19749	0.00134 *
Estimated Covariance Matrix			
v80	v10	v20	v13

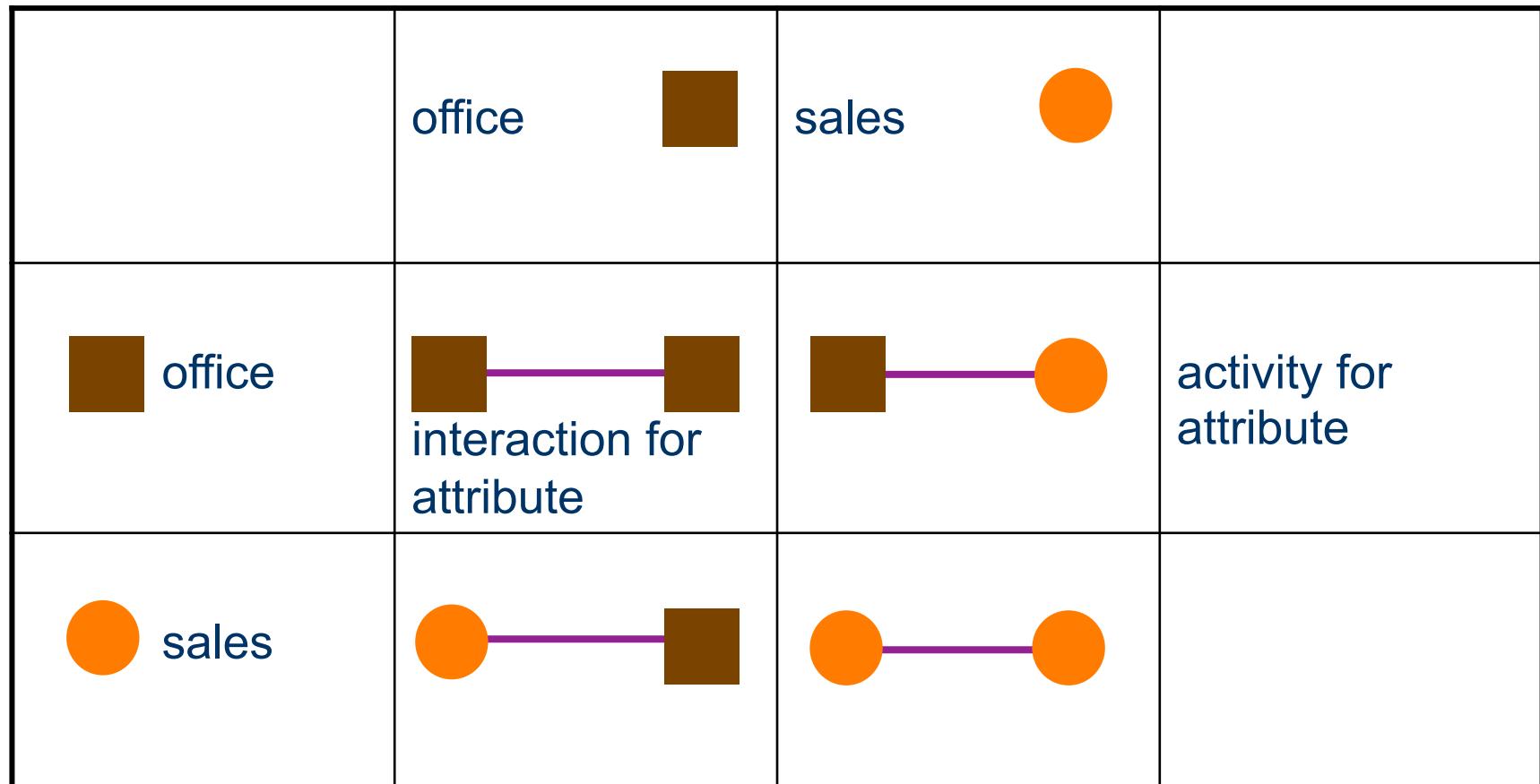
GWESP:  $3t_1(x) - \frac{t_2(x)}{\lambda^1} + \cdots + (-1)^{n-3} \frac{t_{n-2}(x)}{\lambda^{n-3}}$

## Part 4b

# Interpreting attribute-related effects

# Fitting an ERGM in Pnet: a business communications network

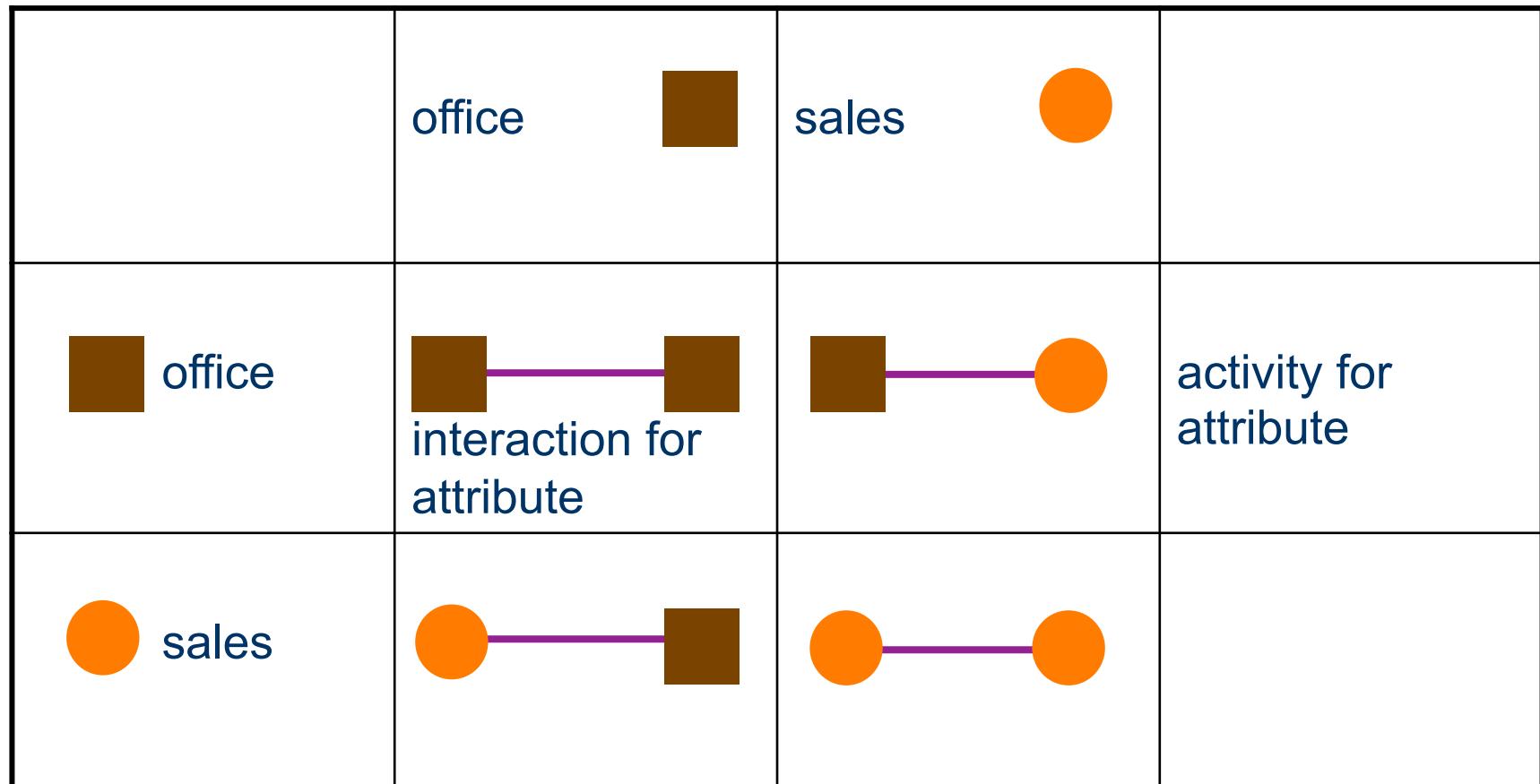
For "wwbusiness.txt" we have recorded whether the employee works in the central office or is a traveling sales representative



# Fitting an ERGM in Pnet: a business communications network

Consider a dyad-independent model

$$\log \Pr(X_{ij} = x_{ij}) = \sigma_1 x_{ij} + \theta_R x_{ij}(OFF_i + OFF_j) + \theta_{Rb} x_{ij} OFF_i OFF_j + \psi(\theta)$$



# Fitting an ERGM in Pnet: a business communications network

With log odds

$$\log \frac{\Pr(X_{ij} = 1)}{\Pr(X_{ij} = 0)} = \sigma_1 + \theta_R (OFF_i + OFF_j) + \theta_{Rb} OFF_i OFF_j$$

$\log \frac{\Pr(X_{ij} = 1)}{\Pr(X_{ij} = 0)}$	office $OFF_j = 1$	sales $OFF_j = 0$	
office $OFF_i = 1$	$\sigma_1 + \theta_R 2 + \theta_{Rb}$ interaction for attribute	$\sigma_1 + \theta_R$	activity for attribute
sales $OFF_i = 0$	$\sigma_1 + \theta_R$	$\sigma_1$	

## Part 4c

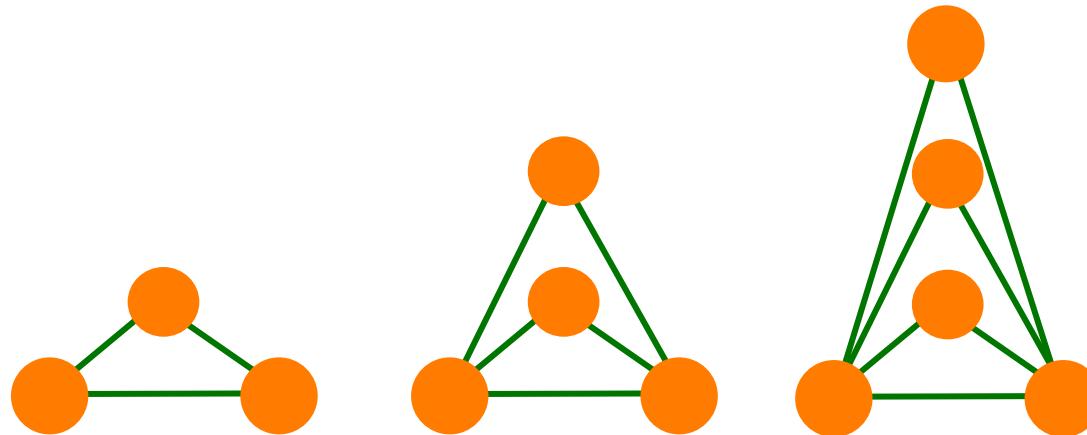
### Interpreting higher order effects

# Unpacking the alternating star effect

Alternating stars a way of

- “**fixing**” the Markov problems (models all degrees)
- **Controlling** for paths in clustering

$$\sigma_2 S_2(x) + \sigma_3 S_3(x) + \cdots + \sigma_{n-1} S_{n-1}(x) = \sigma_{AKS} AKS(x; \lambda)$$



1-triangles:

2-triangles:

3-triangles:

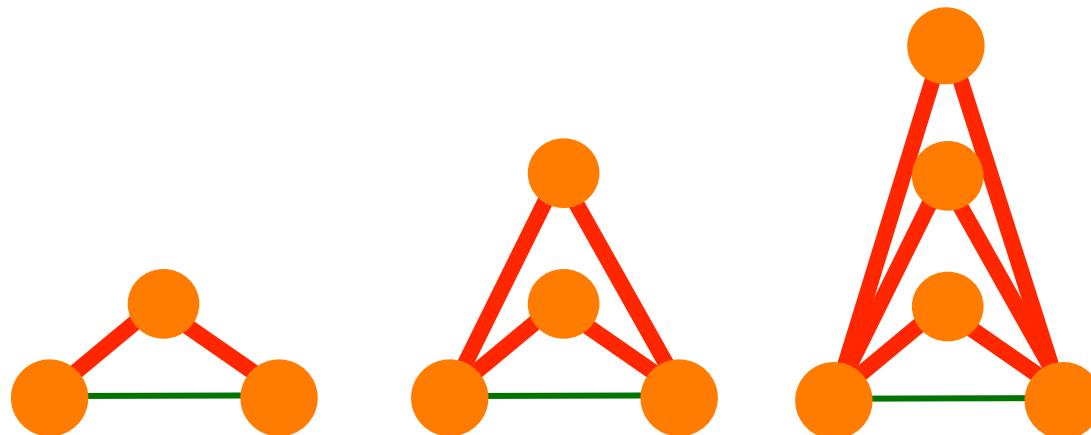
$k$ -triangles measure clustering...

# Unpacking the alternating star effect

Alternating stars a way of

- “fixing” the Markov problems (models all degrees)
- **Controlling** for paths in clustering

$$\sigma_2 S_2(x) + \sigma_3 S_3(x) + \cdots + \sigma_{n-1} S_{n-1}(x) = \sigma_{AKS} AKS(x; \lambda)$$

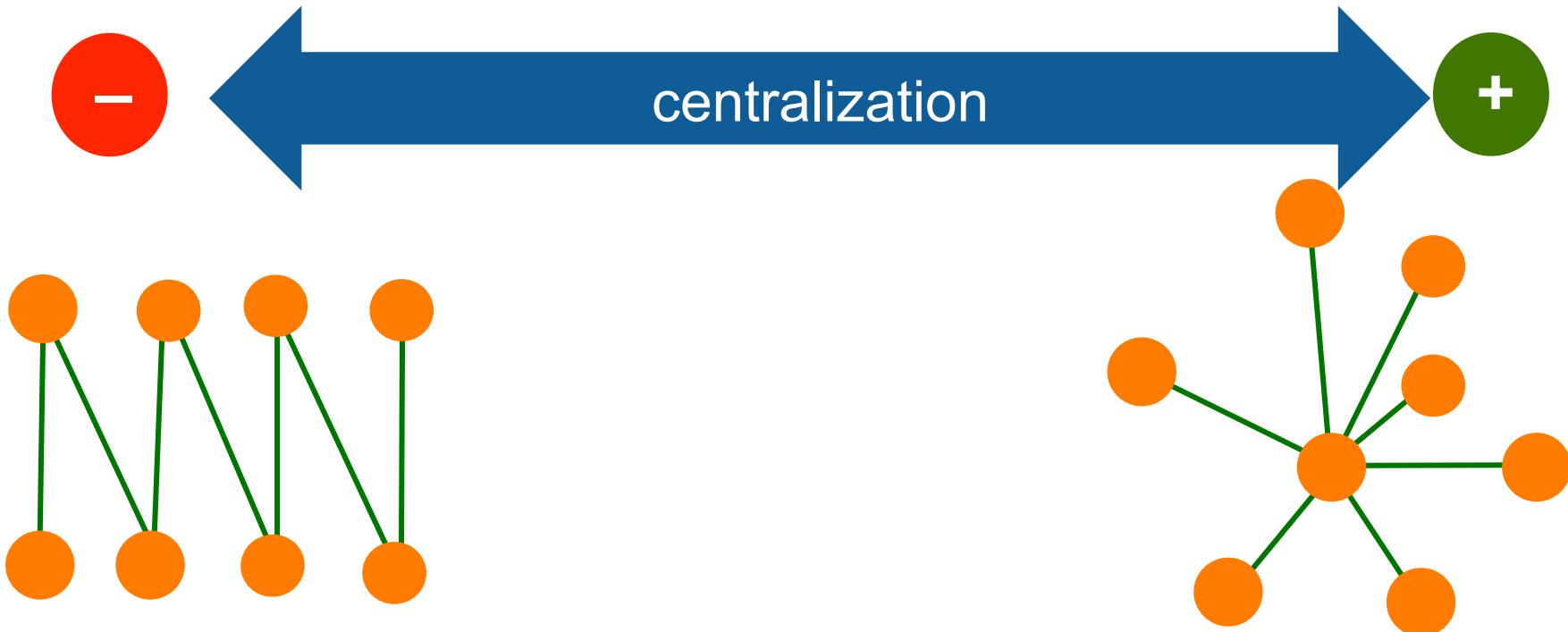


1-triangles:    2-triangles:    3-triangles:

Is it **closure** or an **artefact** of many stars/2-paths?

# Unpacking the alternating star effect

Interpreting the alternating star **parameter**:



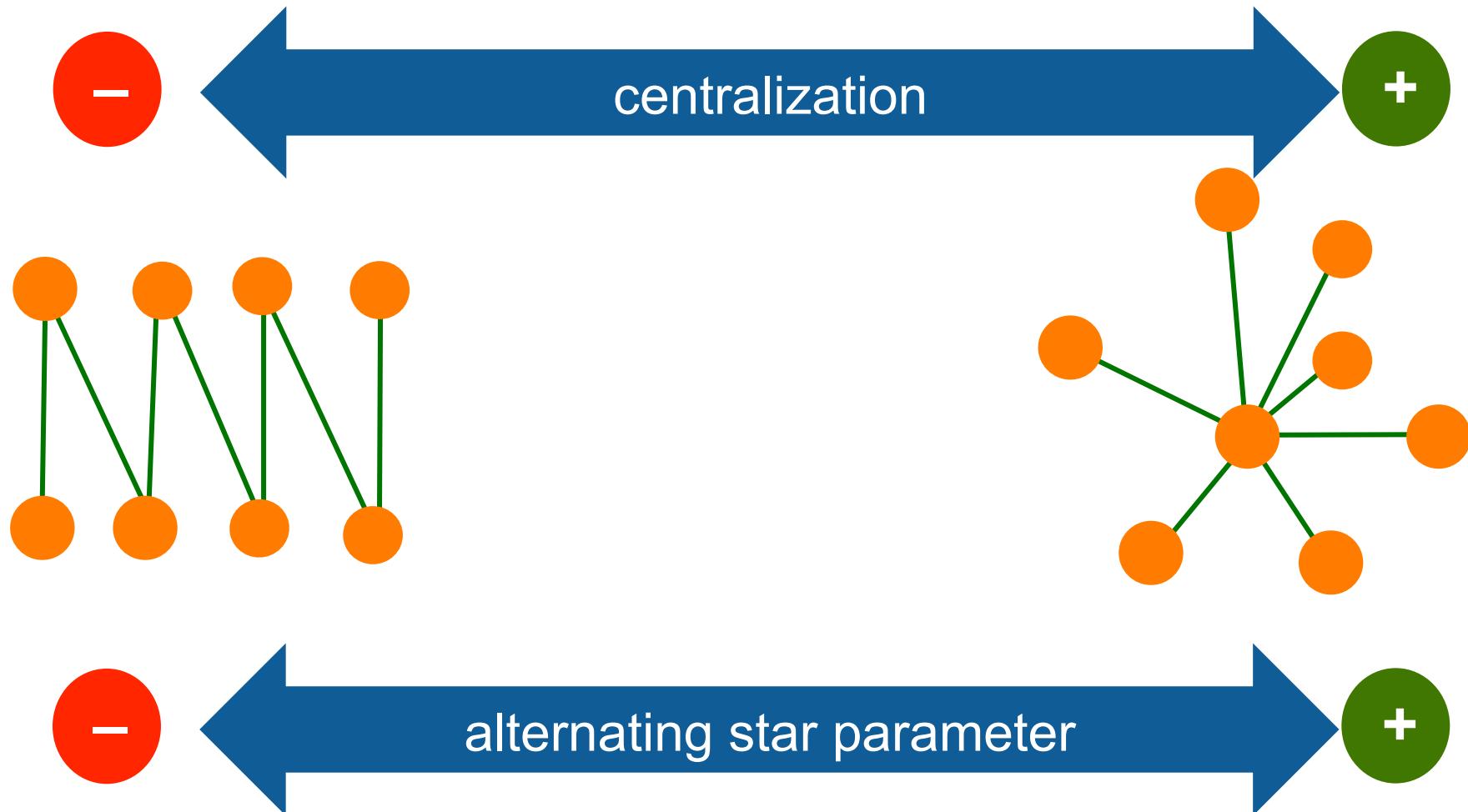
$$Var(x_{i+}) = \sum_i \frac{(x_{i+} - \bar{x})^2}{n-1} = .21$$

$$Var(x_{i+}) = 4.42$$

variance of degree measure centralization

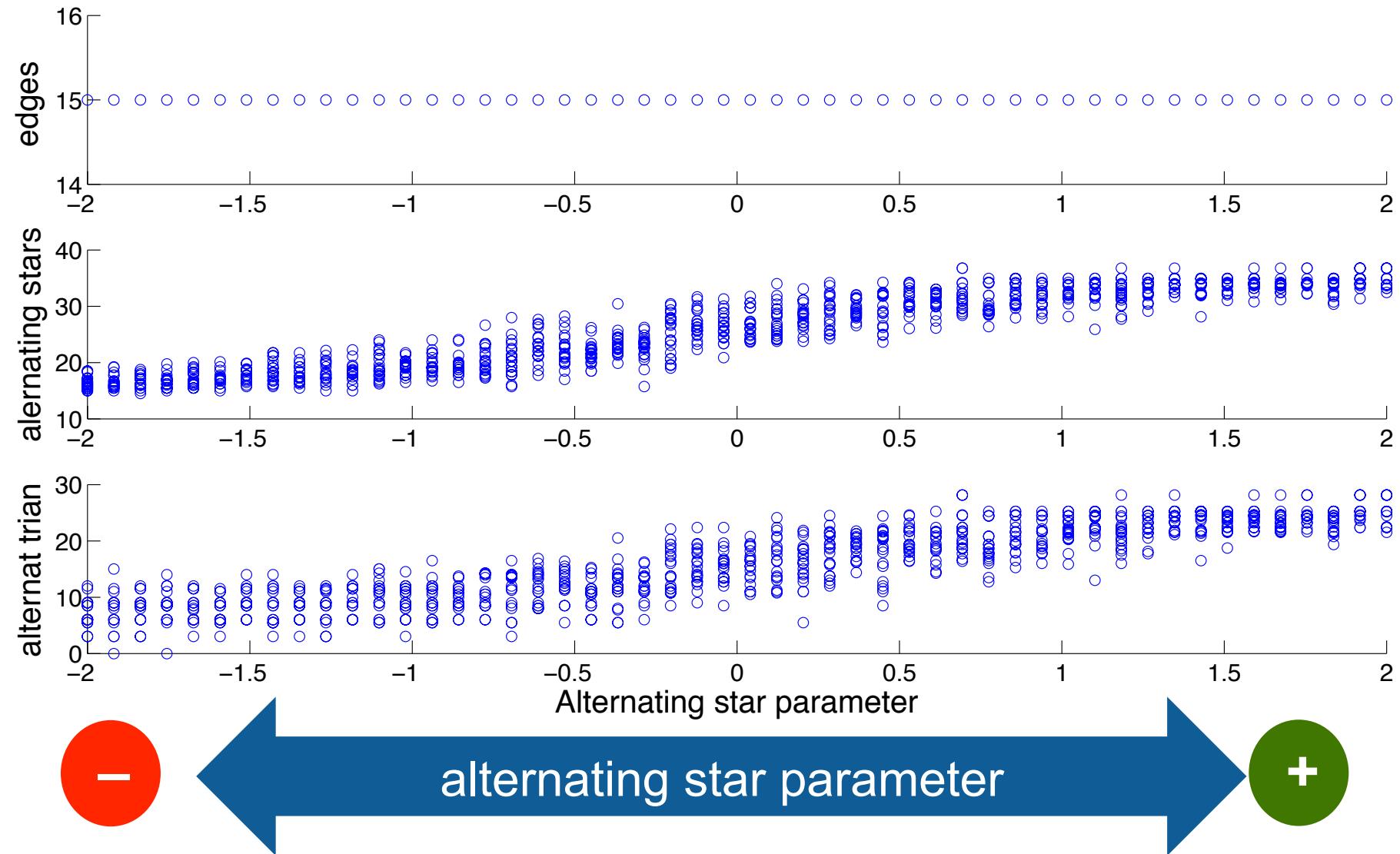
# Unpacking the alternating star effect

Interpreting the alternating star **parameter**:



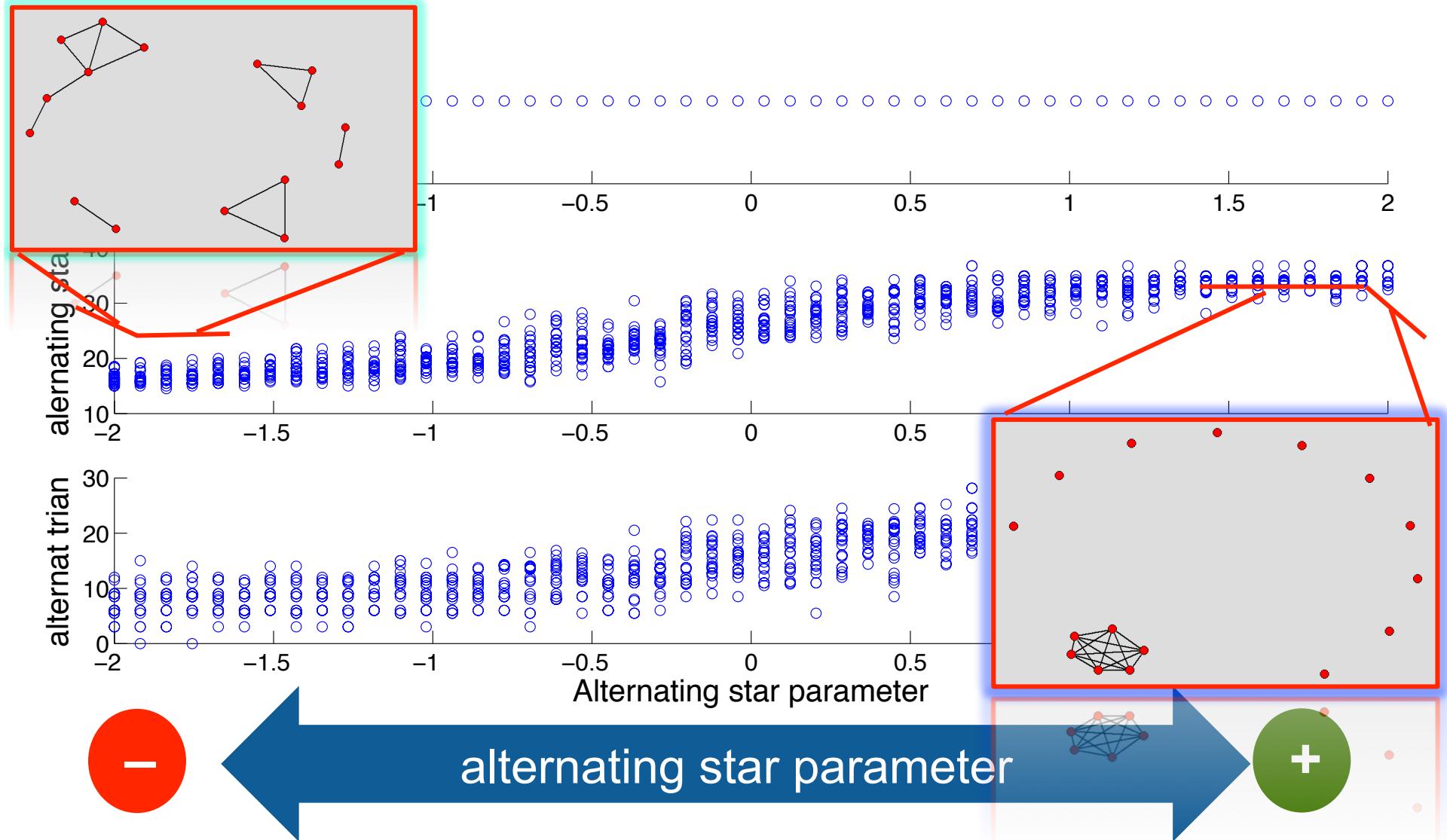
# Unpacking the alternating star effect

Statistics for graph ( $n = 16$ ); fixed density; alt trian: 1.17



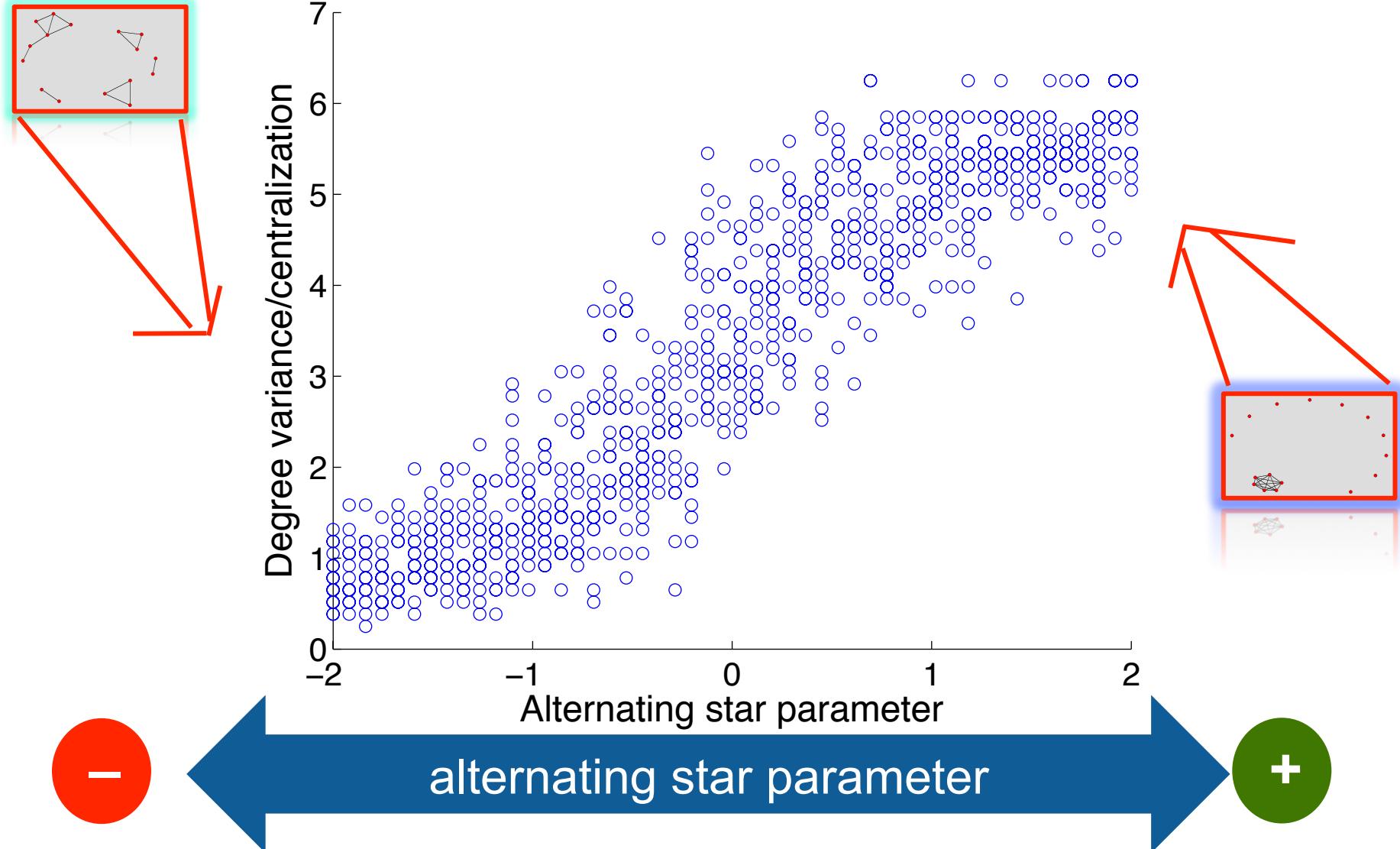
# Unpacking the alternating star effect

Statistics for graph ( $n = 16$ ); fixed density; alt trian: 1.17



# Unpacking the alternating star effect

Graphs ( $n = 16$ ); fixed density; alt trian: 1.17



## Problem with Markov models

Note also the influence of isolates:

Alt, stars in terms  
of degree distribution

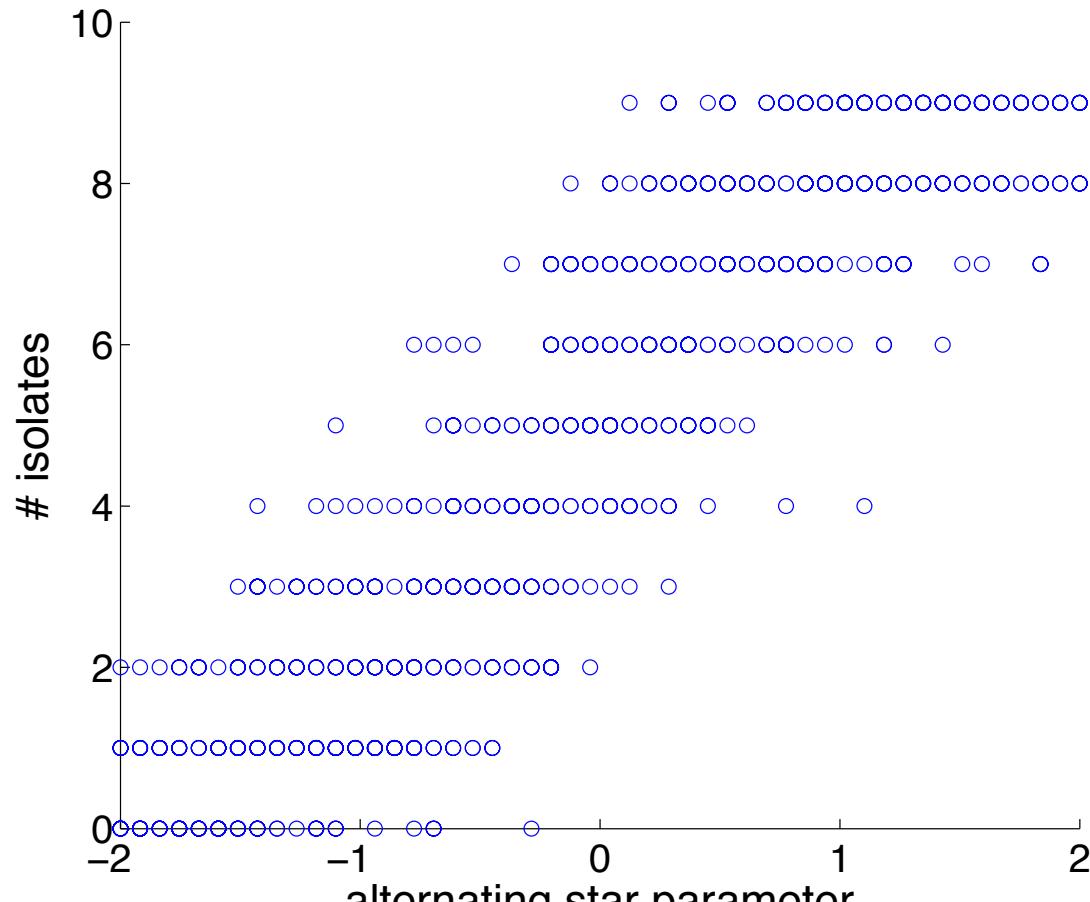
$$= \left( \frac{1}{1 - e^{-\alpha}} \right)^2 \sum_{j=0}^{n-1} d_j(x) e^{-\alpha j} + \frac{2L(x)}{1 - e^{-\alpha}} - \frac{n}{(1 - e^{-\alpha})^2}$$

$$d_j(x) = \#\{i : x_{i+} = j\}$$

$$\lambda = e^\alpha / (e^\alpha - 1)$$

# Problem with Markov models

Note also the influence of isolates:



-

+

alternating star parameter

## Problem with Markov models

Note also the influence of isolates:

This is because we impose a particular shape on the degree distribution

$$\begin{aligned}\sigma_2 S_2(x) + \sigma_3 S_3(x) + \cdots + \sigma_{n-1} S_{n-1}(x) &= \sigma_{AKS} AKS(x; \lambda) \\ &= \left( \frac{1}{1 - e^{-\alpha}} \right)^2 \sum_{j=0}^{n-1} d_j(x) e^{-\alpha j} + \frac{2L(x)}{1 - e^{-\alpha}} - \frac{n}{(1 - e^{-\alpha})^2}\end{aligned}$$

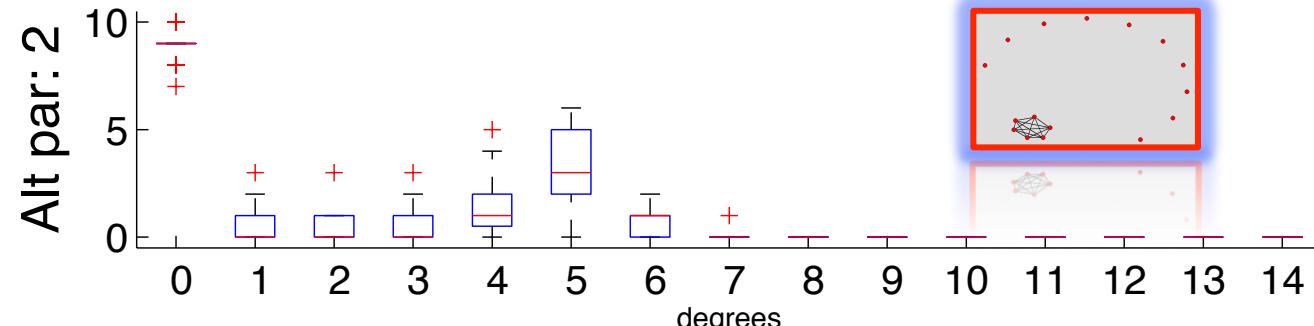
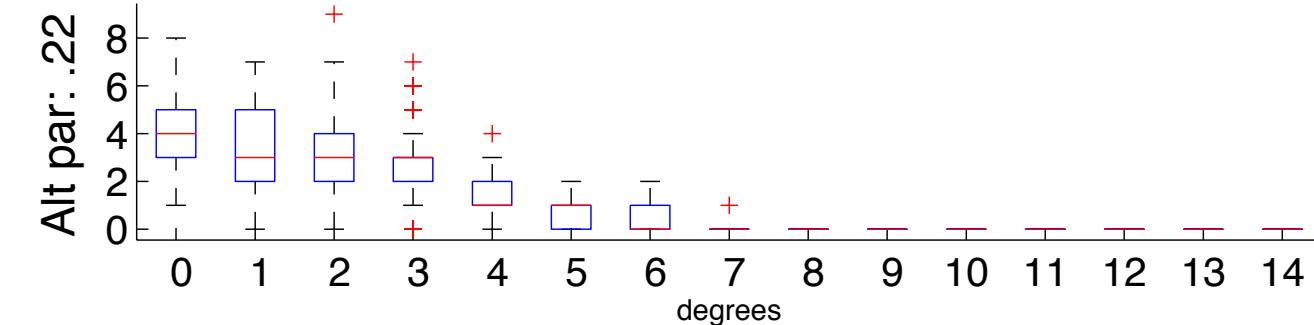
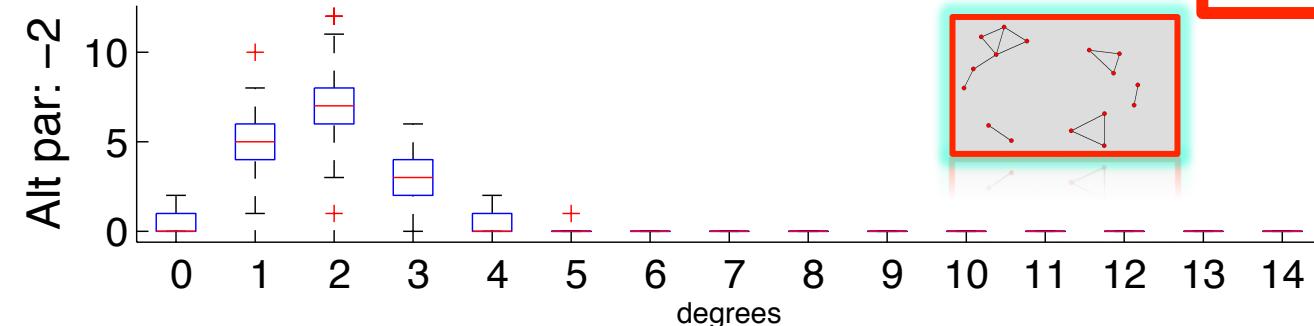
# Problem with Markov models

The degree distribution for different values of:

$\sigma_{AKS}$

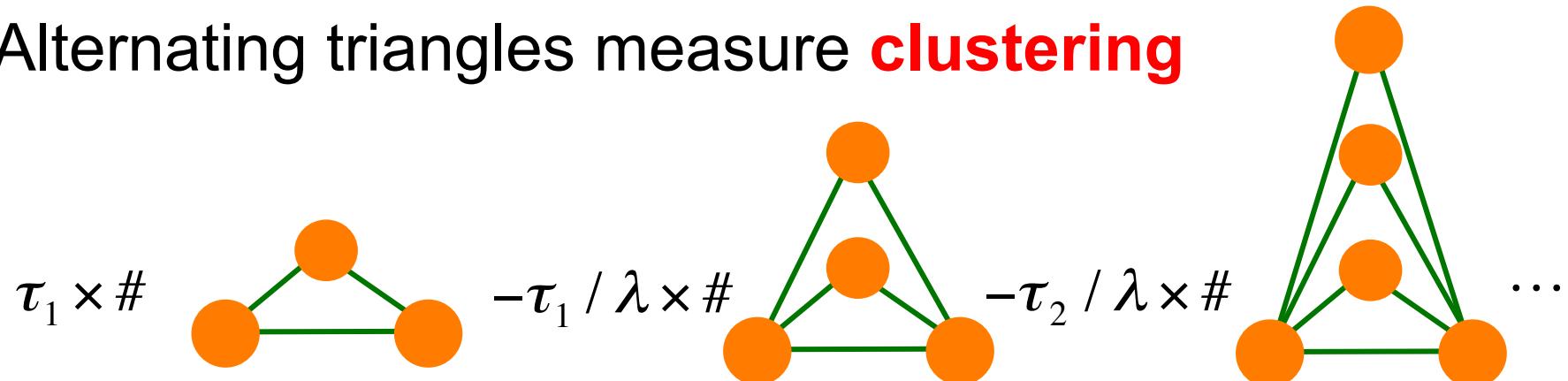


alterning star parameter



# Unpacking the alternating triangle effect

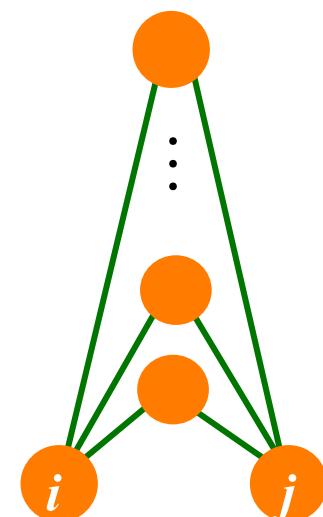
Alternating triangles measure **clustering**



We may also define the  
**Edgewise Shared Partner Statistic:**

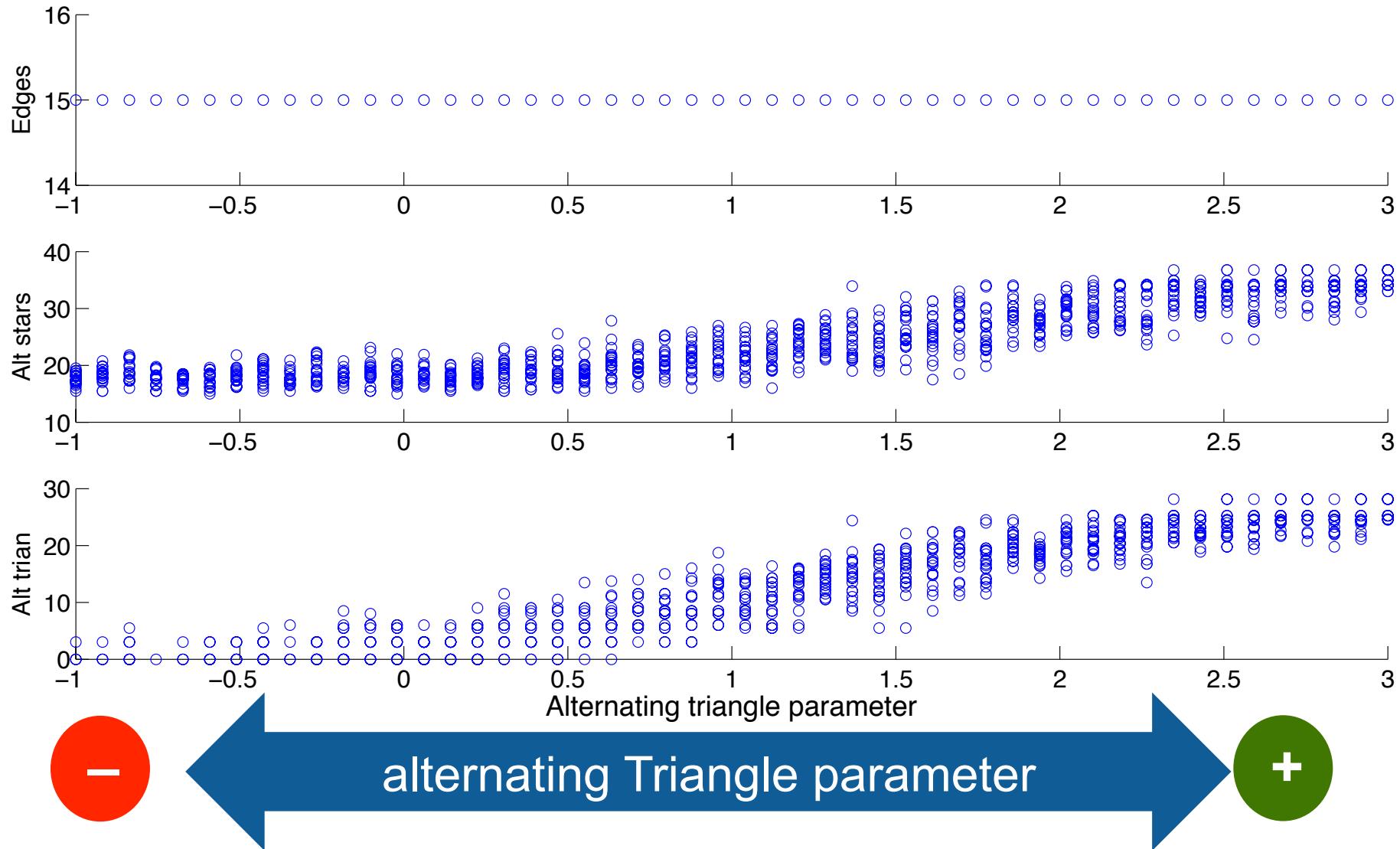
$$ESP_k = \#\{(i, j) : i \sim j, S_{2ij} = k\}$$

... and we can weigh together the **ESP** statistics using **Geometrically decreasing weights: GWESP**



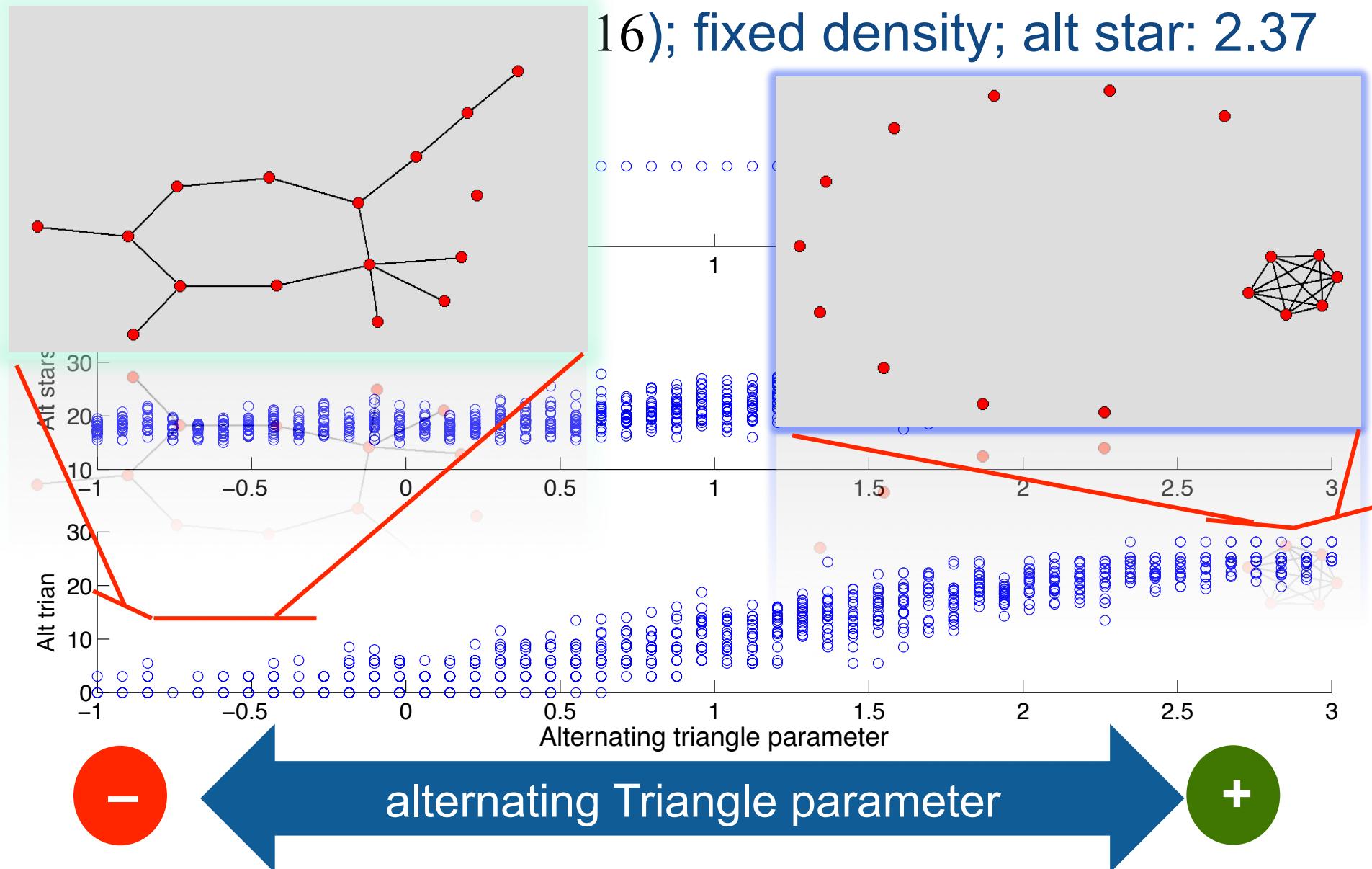
# Unpacking the alternating triangle effect

Statistics for graph ( $n = 16$ ); fixed density; alt star: 2.37



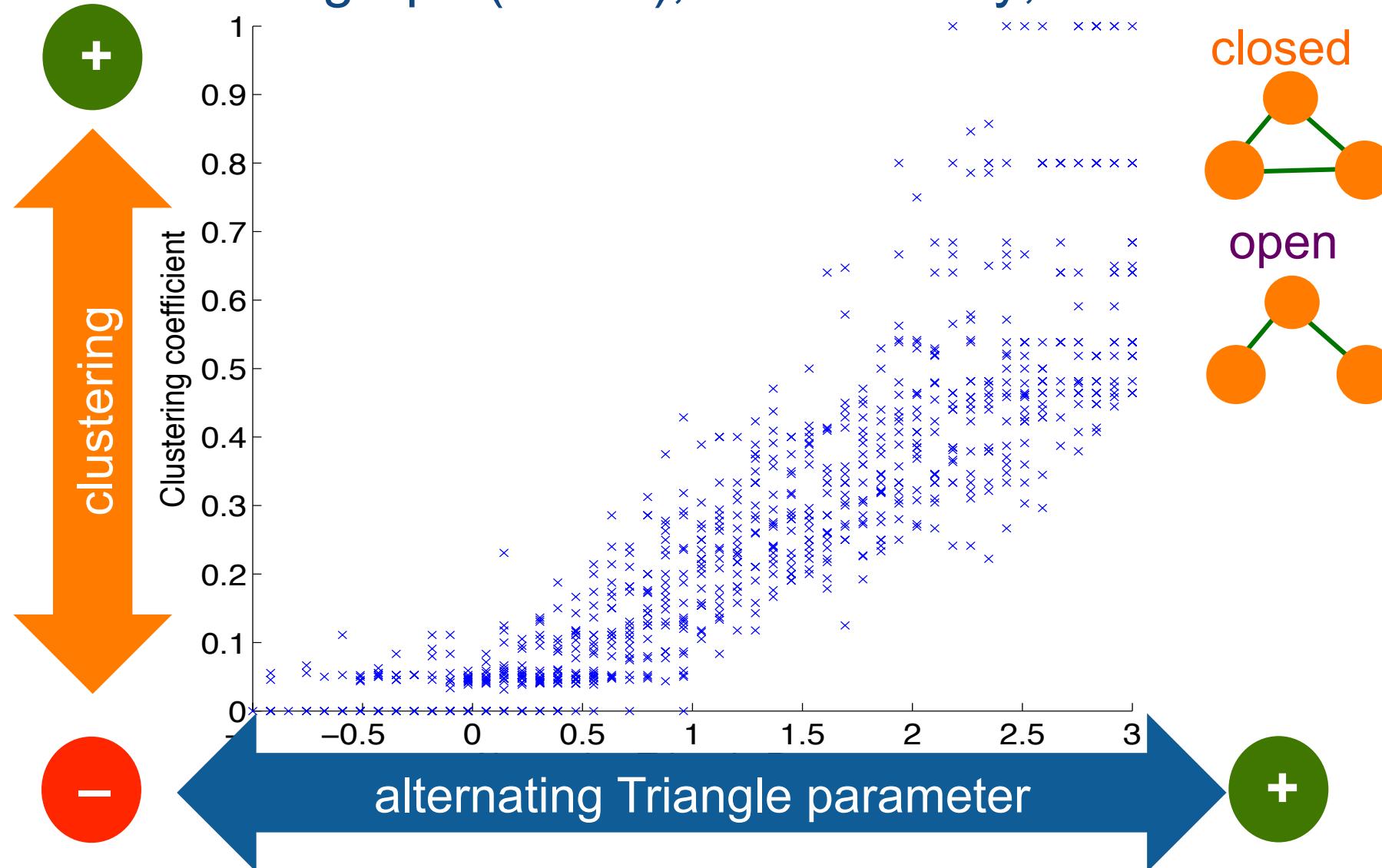
# Unpacking the alternating triangle effect

16); fixed density; alt star: 2.37



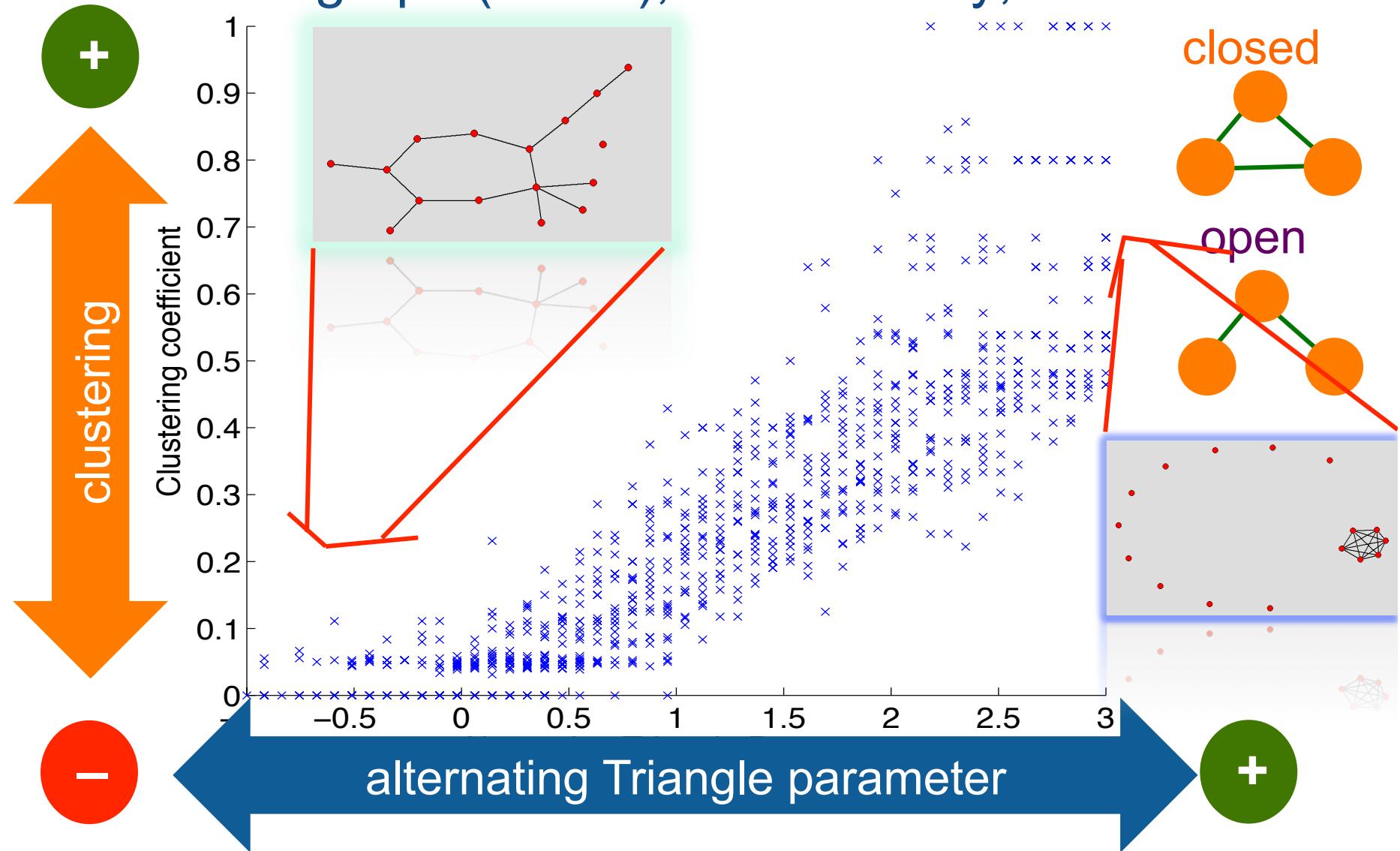
# Unpacking the alternating triangle effect

Statistics for graph ( $n = 16$ ); fixed density; alt star: 2.37



# Unpacking the alternating triangle effect

Statistics for graph ( $n = 16$ ); fixed density; alt star: 2.37



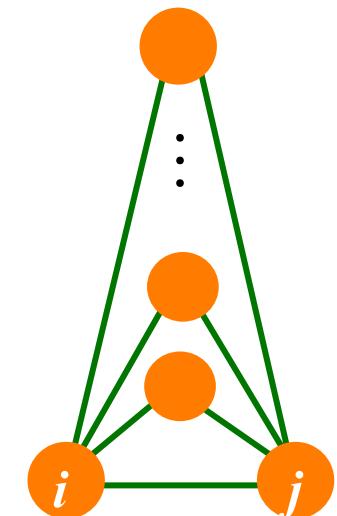
# Unpacking the alternating triangle effect

Alternating triangles model **multiply clustered areas**

For multiply clustered areas triangles stick together  
We model how many others tied actors have

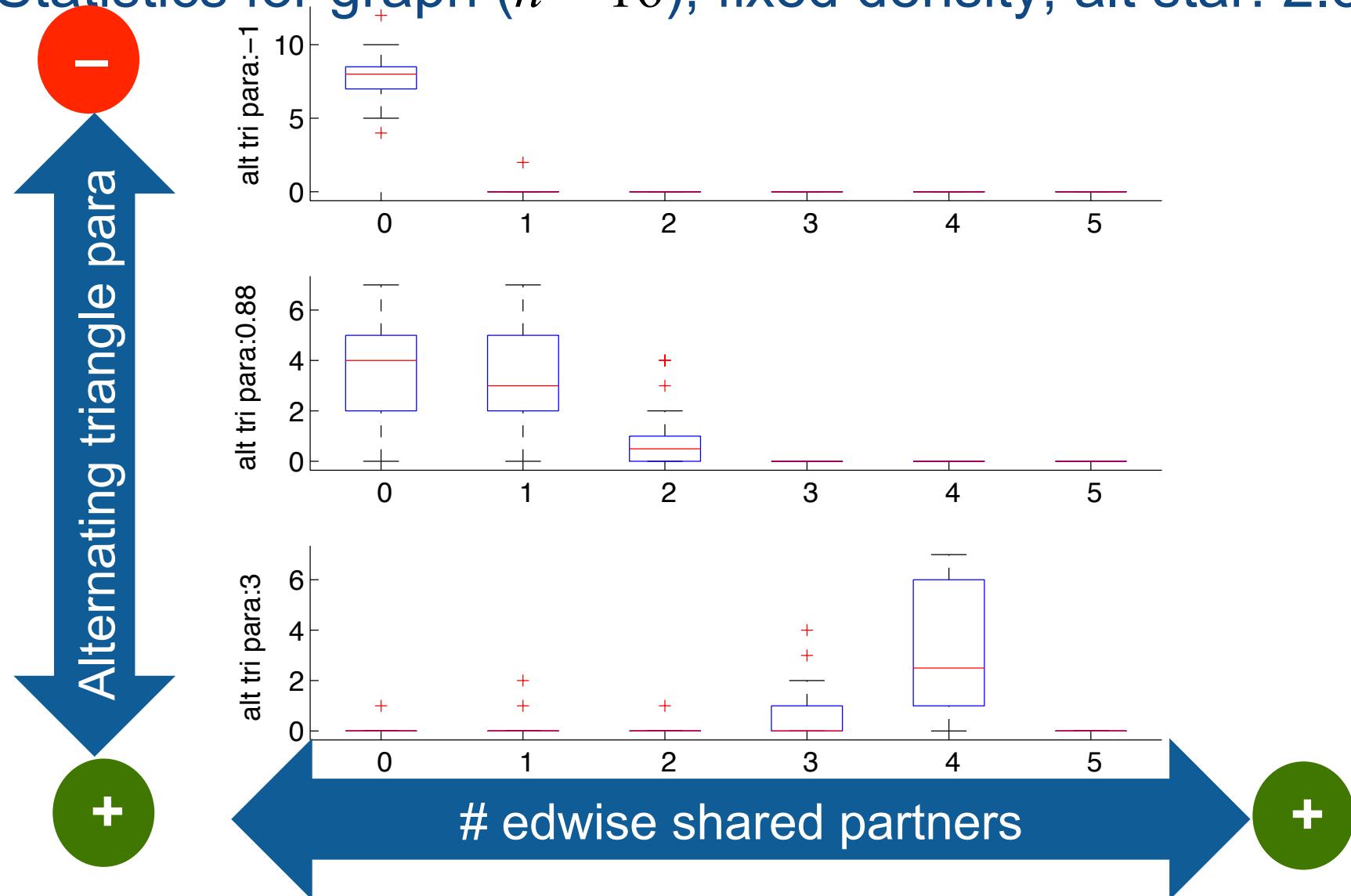
**Edgewise Shared Partner Statistic:**

$$ESP_k = \#\{(i, j) : i \sim j, S_{2ij} = k\}$$



# Unpacking the alternating triangle effect

Statistics for graph ( $n = 16$ ); fixed density; alt star: 2.37



## Part 5

# Convergence check and goodness of fit

# Revisiting the Florentine families



Formula: BusyNetNet ~ kstar(1:3) + triangles

Newton-Raphson iterations: 42

MCMC sample of size 10000

**statnet**

Monte Carlo MLE Results:

	Estimate	Std. Error	MCMC s.e.	p-value
kstar1	-1.6130	0.6699	0.462	0.0176 *
kstar2	0.7492	0.6407	0.455	0.2446
kstar3	-0.5408	0.3574	0.225	0.1330
triangle	1.483	0.4550	0.33	0.0111 **
---				

Why difference?

estimation-padgetestsunday.txt - Notepad

PNet

```
* num of iterations in each step = 280.000000
*****
mean statistics in phase3:15.246000 37.306000 25.856000 5.054000

*Estimation Result for Network SUMMARY (parameter, standard error, t-statistics)
NOTE: t-statistics = (observation - sample mean)/standard error
      Edge: -4.137319, 1.07210, -0.04370 *
      2-Star: 0.973517, 0.59180, -0.06408
      3-Star: -0.563624, 0.35420, -0.09745
      Triangle: 1.261550, 0.61588, -0.01524 *
```

# Revisiting the Florentine families

$$L(x) = \# \text{ } \begin{array}{c} \text{---} \\ | \\ \text{---} \end{array} \quad S_2(x) = \# \text{ } \begin{array}{c} \text{---} \\ | \\ \text{---} \\ | \\ \text{---} \end{array} \quad S_3(x) = \# \text{ } \begin{array}{c} \text{---} \\ | \\ \text{---} \\ | \\ \text{---} \\ | \\ \text{---} \end{array} \quad T(x) = \# \text{ } \begin{array}{c} \text{---} \\ | \\ \text{---} \\ | \\ \text{---} \end{array}$$

## Pnet checks convergence

$$E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$$

in 3rd phase

```
estimation-padgetestsunday.txt - Notepad
File Edit Format View Help
* num of iterations in each step = 280.000000
*****
mean statistics in phase3:15.246000 37.306000 25.856000 5.054000
*Estimation Result for Network SUMMARY (parameter, standard error, t-statistics)
NOTE: t-statistics = (observation - sample mean)/standard error
      Edge: -4.137319, 1.07210, -0.04370 *
      2-Star: 0.973517, 0.59180, -0.06408
      3-Star: -0.563624, 0.35420, -0.09745
      Triangle: 1.261550, 0.61588, -0.01524 *
```

# Revisiting the Florentine families



Formula: `BusyNetNet ~ kstar(1:3) + triangles`

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MCMC sample of size 10000

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	Estimate	Std. Error	MCMC s.e.	p-value
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kstar2	0.7492	0.6407	0.455	0.2446
kstar3	-0.5408	0.3574	0.225	0.1330
triangle	1.4837	0.4592	0.138	0.0016 **
---				
Signif. codes:	0 ****	0.001 ***	0.01 **	0.05 *
	.	.	0.1	' '
				1

Lets check  
For statnet

$$E_{\hat{\theta}_{MLE}} \{z(X)\} = z(x_{obs})$$

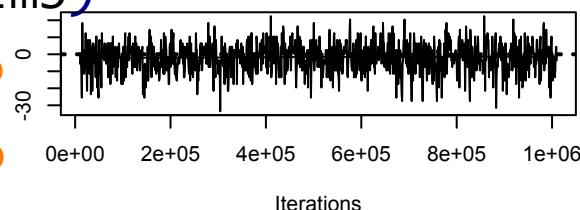
```
Estim3 <- ergm(BusyNetNet ~ kstar(1:3) + triangles ,  
verbose=TRUE)
```

```
mcmc.diagnostics(Estim3)
```

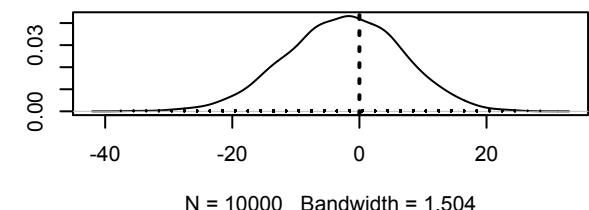
$$L(x) = \#$$

Summary of MCMC samples

Trace of kstar1

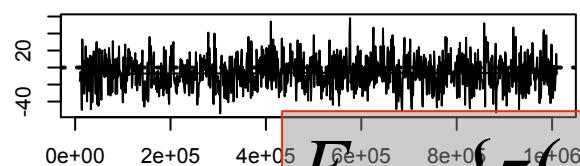


Density of kstar1

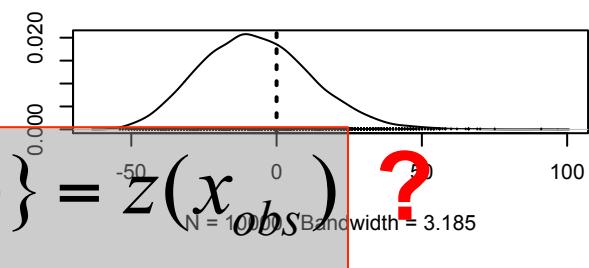


$$S_2(x) = \#$$

Trace of kstar2

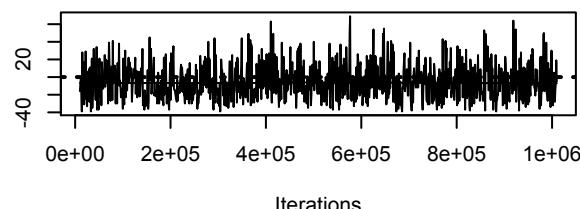


Density of kstar2

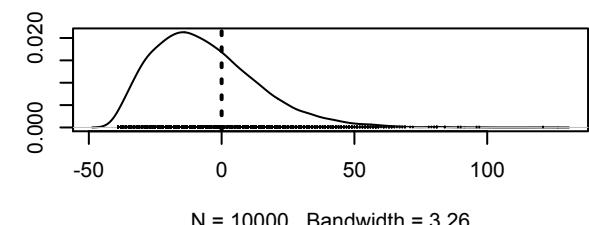


$$S_3(x) = \#$$

Trace of kstar3

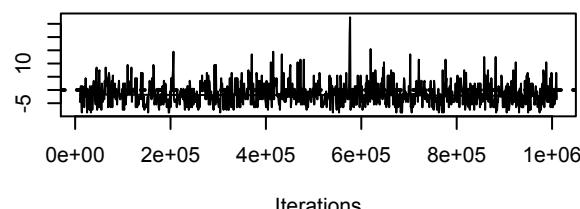


Density of kstar3

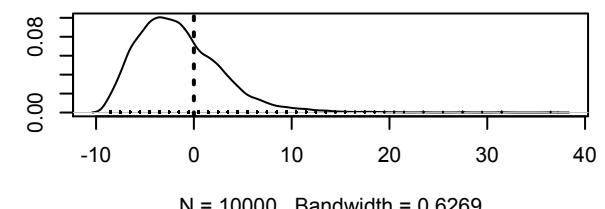


$$T(x) = \#$$

Trace of triangle



Density of triangle



Given a specific model we can simulate potential outcomes under that model. This is used for

- Estimation:
  - (a) to match observed statistics and expected statistics
  - (b) to check that we have “the solution”
- GOF: to check whether the model can replicated features of the data that we not explicitly modelled
- Investigate behaviour of model, e.g.: degeneracy and dependence on scale

## Simulation, GOF, and Problems

Standard goodness of fit procedures are not valid for ERGMs – no F-tests or Chi-square tests available

If indeed the fitted model adequately describes the data generating process, then the graphs that the model produces should be similar to observed data

For fitted effects this is true by construction

For effects/structural feature that are not fitted this may not be true

If it is true, the modelled effects are the only effects necessary to produce data – a “proof” of the concept or ERGMs

## Example: for our fitted model for Lazega

```
NOTE: t-statistics = (observation - sample mean)/standard error
      effects           estimates      stderr   t-ratio
      edge              -5.862515    0.56404  0.04105 *
      AT(2.00)          1.011721    0.17095  0.05003 *
      practice_interaction 1.499409    0.40322  0.02371 *
      practice_activity   -0.331023   0.21995  0.02142
      senior_sum          0.842661    0.23348  0.04943 *
      sex_matching         0.702477   0.26389  0.05839 *
      off_matching         1.145290    0.19749  0.00134 *
```

We can look at how well the model reproduces

$$g(x_{obs})$$

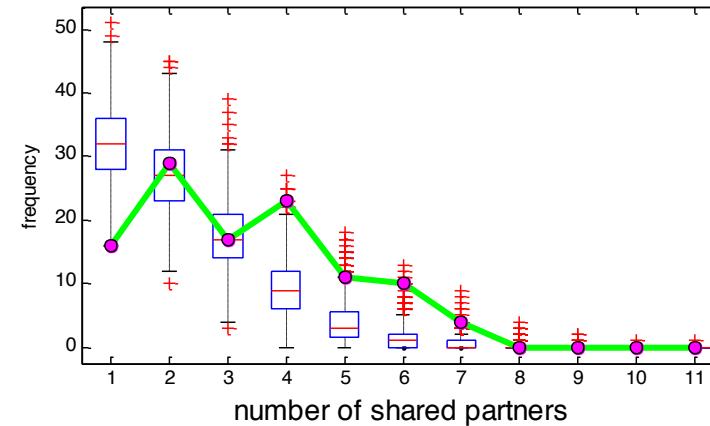
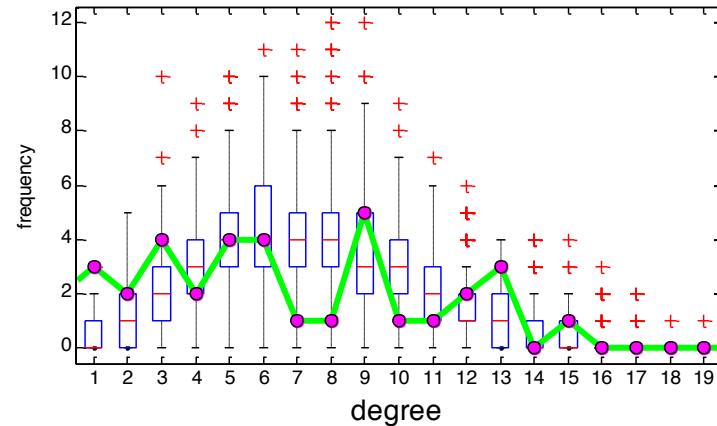
For arbitrary function  $g$

# Simulation, GOF, and Problems

From the goodness-of-fit tab in Pnet we get

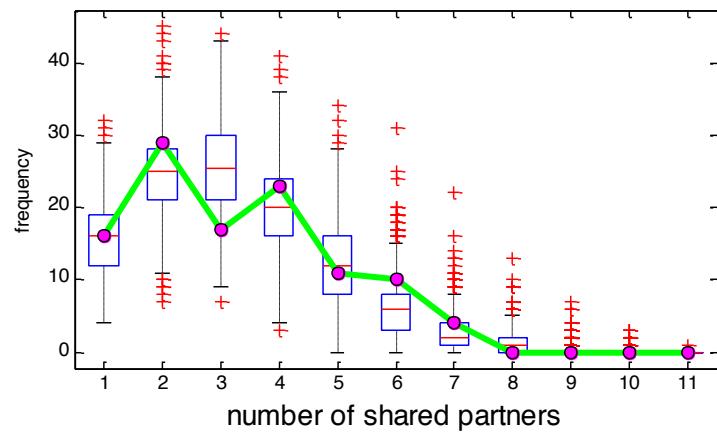
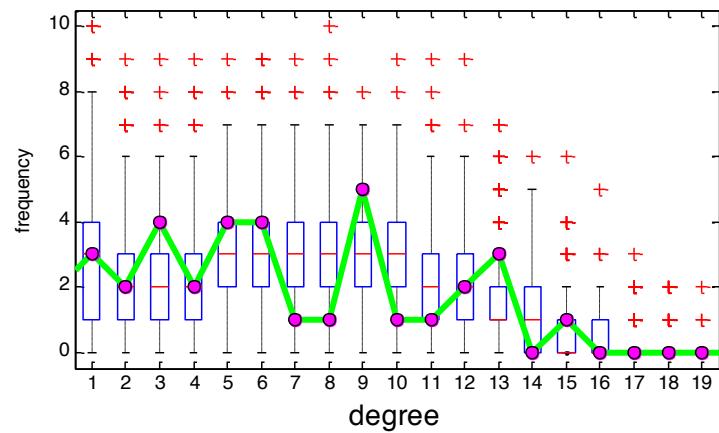
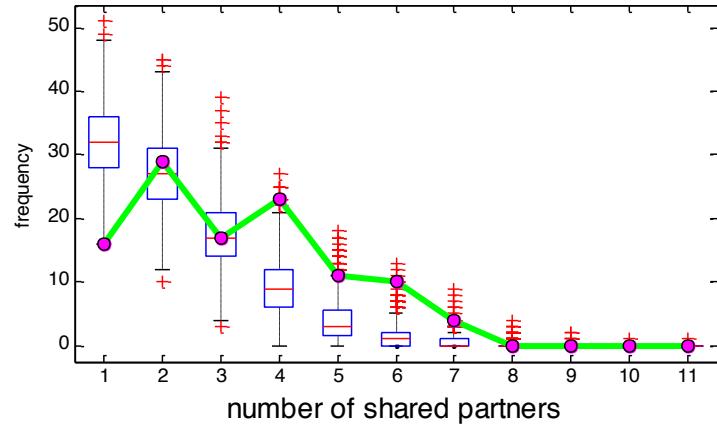
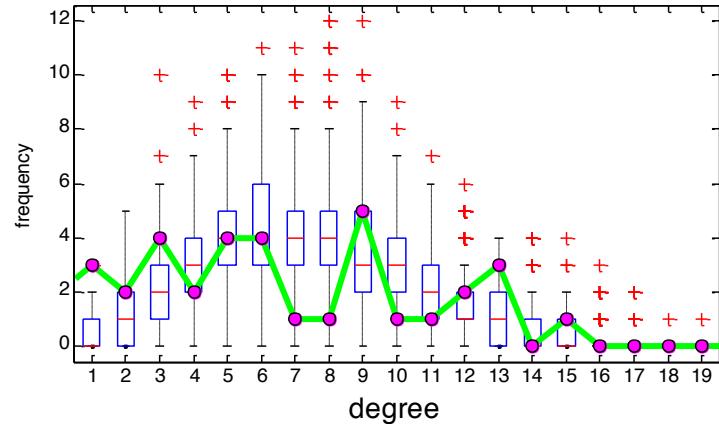
<b>Effect</b>	<b>MLE</b>	<b>s.e.</b>
Density	-6.501	0.727
Main effect seniority	1.594	0.324
Main effect practice	0.902	0.163
Homophily effect practice	0.879	0.231
Homophily sex	1.129	0.349
Homophily office	1.654	0.254

# Simulation, GOF, and Problems



Effect	MLE	s.e.	MLE	s.e.
Density	-6.501	0.727	-6.510	0.637
Main effect seniority	1.594	0.324	0.855	0.235
Main effect practice	0.902	0.163	0.410	0.118
Homophily effect practice	0.879	0.231	0.759	0.194
Homophily sex	1.129	0.349	0.704	0.254
Homophily office	1.654	0.254	1.146	0.195
GWEPS			0.897	0.304
Log-lambda			0.778	0.215

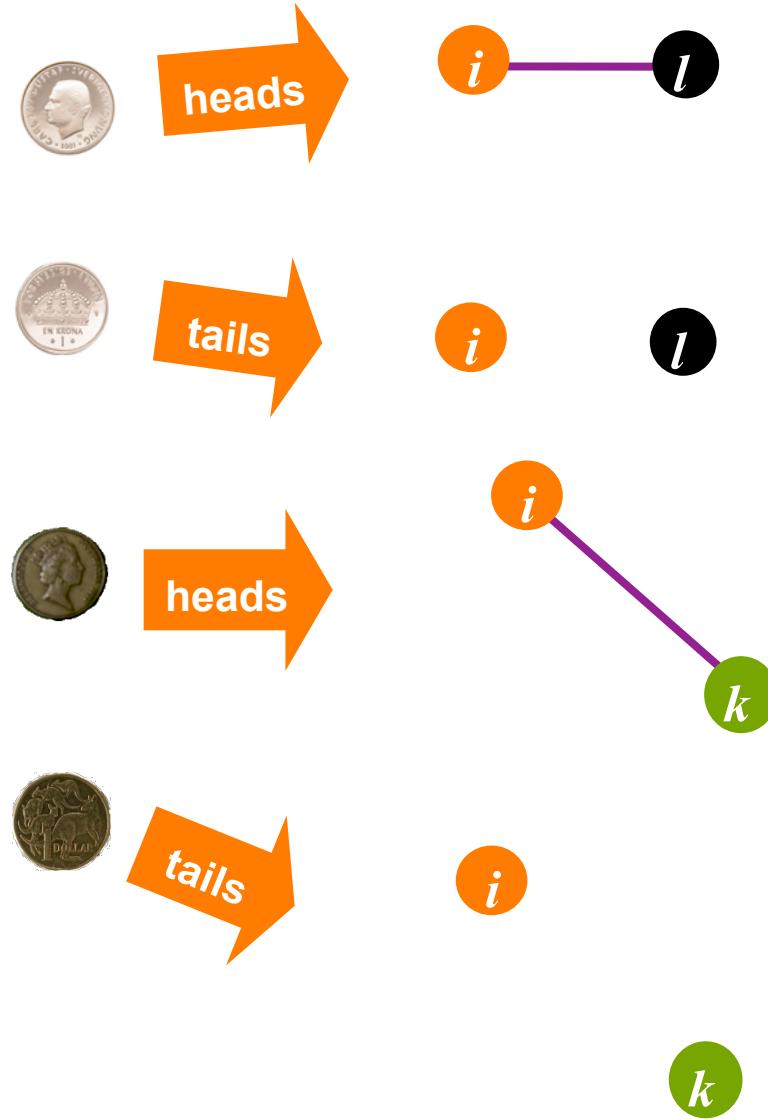
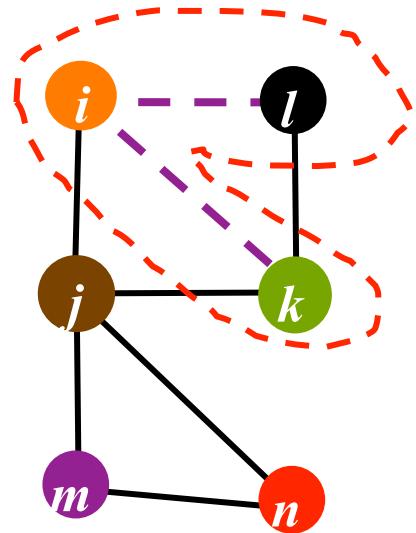
# Simulation, GOF, and Problems



## Part 6

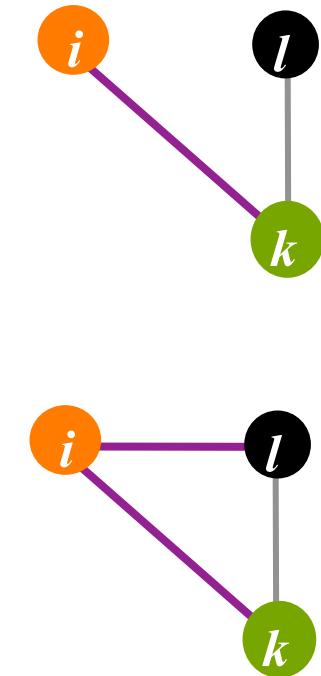
Dependencies – Sufficient statistics -  
homogeneity

# Independence - Deriving the ERGM



# Independence - Deriving the ERGM

AUD	
	0.5
	0.5
SEK	
	0.5
	0.25
	0.25
	0.25
	0.25



Knowledge of AUD, e.g. does **not** help us predict SEK  
e.g. whether or

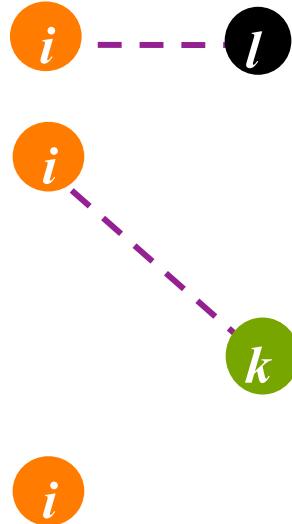
# Independence - Deriving the ERGM

Knowledge of AUD, e.g.  does **not** help us predict SEK

e.g. whether  or 

even though **dyad**  $\{i, l\}$  

and **dyad**  $\{i, k\}$



have **vertex**  $i$

in **common**

# Independence - Deriving the ERGM

May we find model such that knowledge of AUD,  
e.g. **does** help us predict SEK

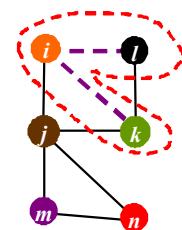
e.g. whether  or  ?

SEK		0.5		0.5

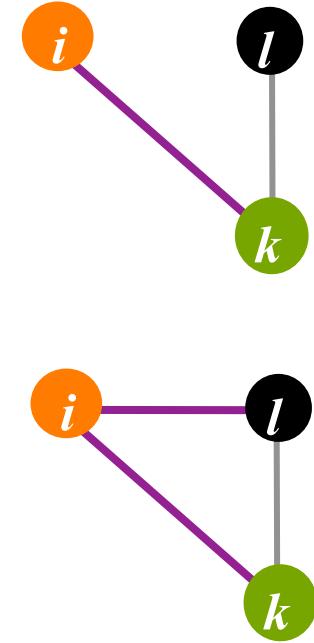
AUD	
	0.5
	0.5
	0.4
	0.1
	0.1
	0.4



?

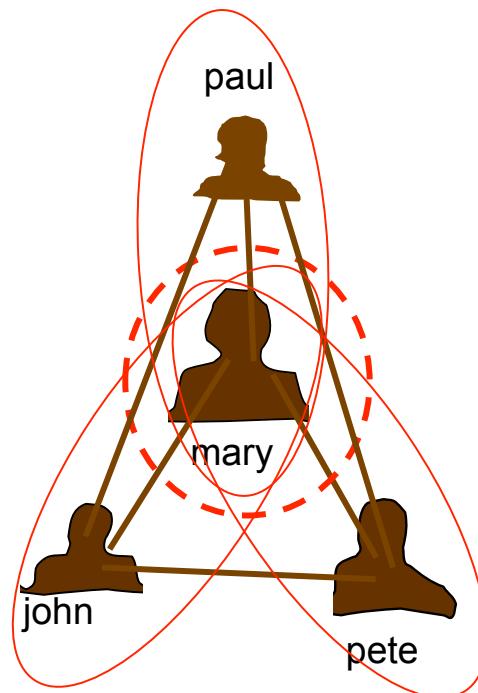


AUD



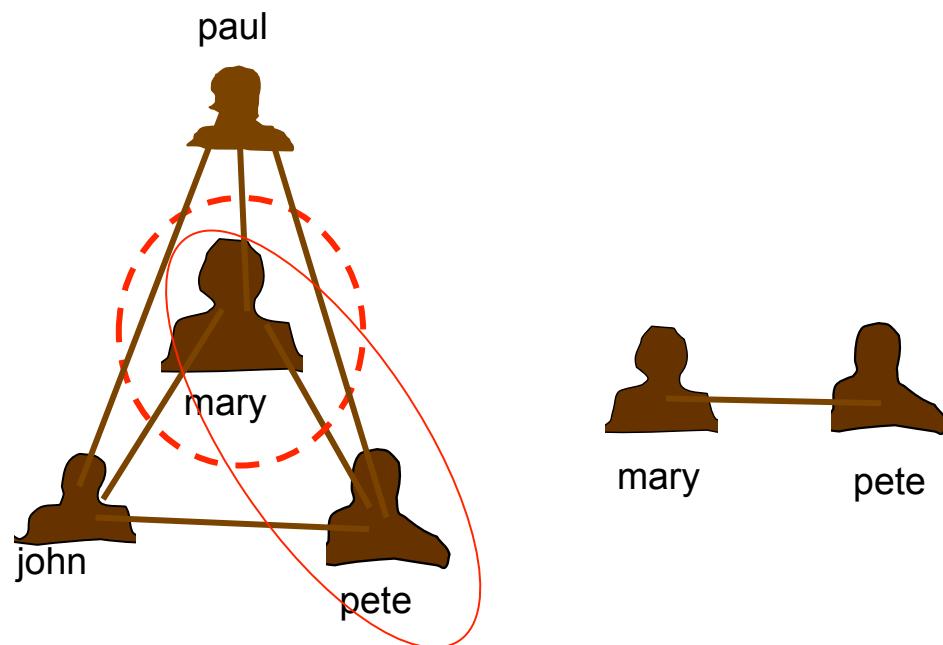
# Deriving the ERGM: From Markov graph to Dependence graph

Consider the tie-variables that have **Mary** in  
**common**

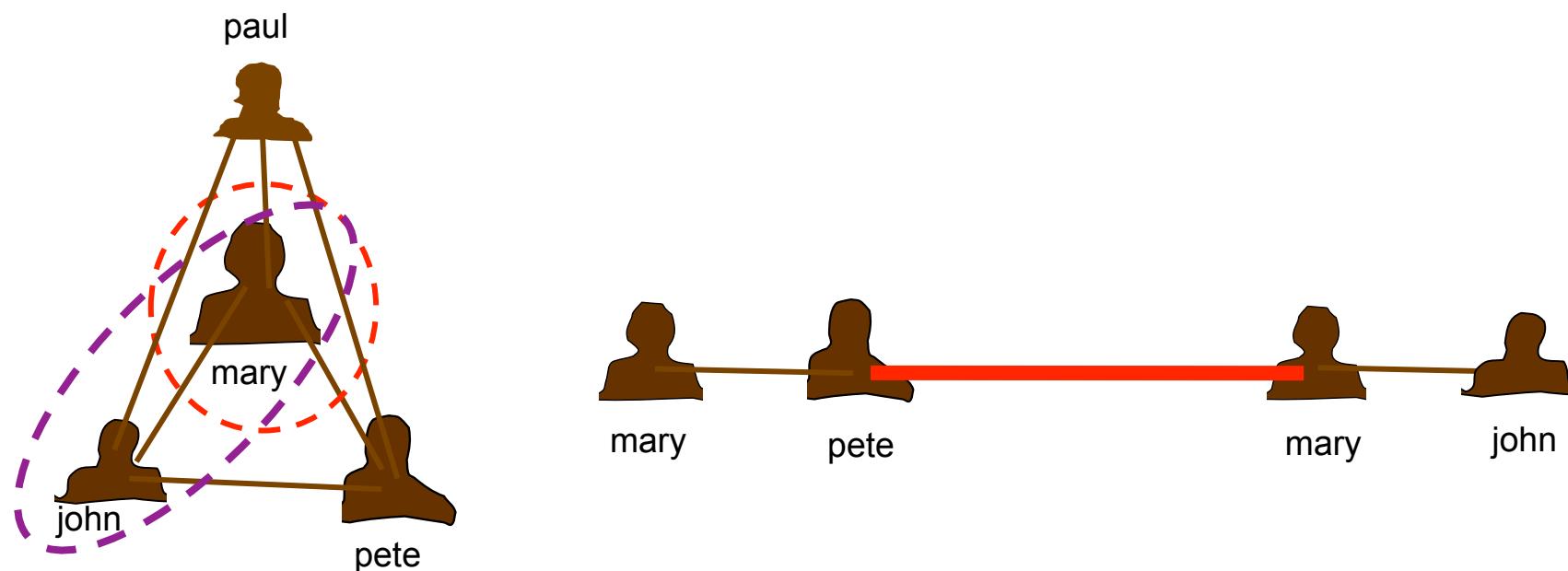


How may we make these “**dependent**”?

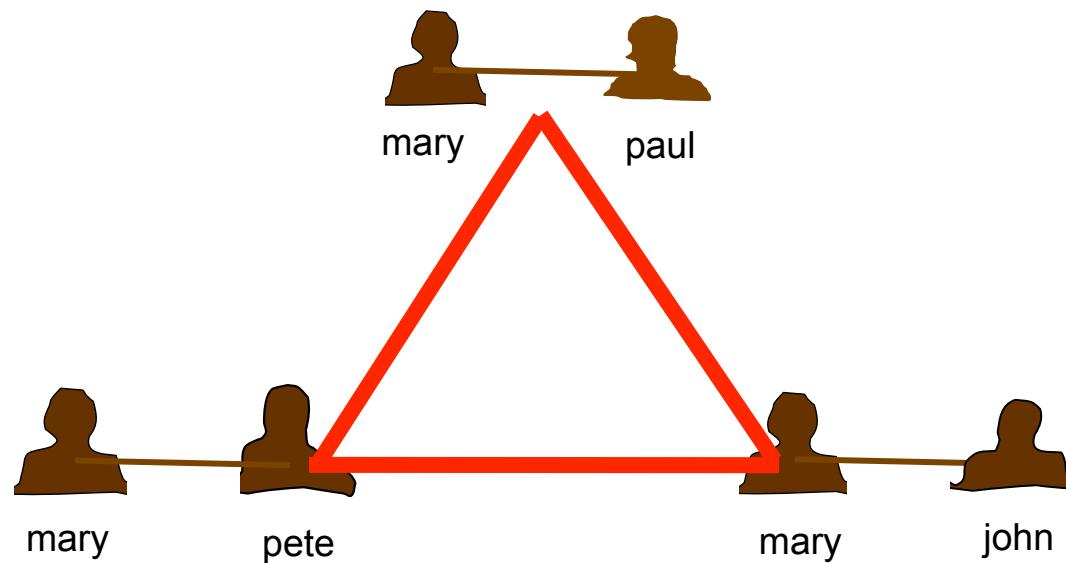
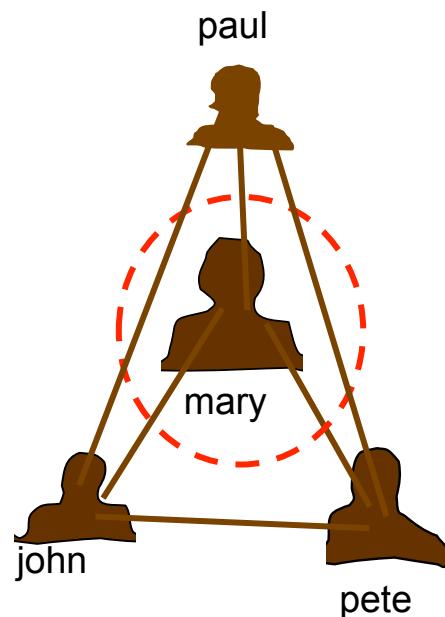
# Deriving the ERGM: From Markov graph to Dependence graph



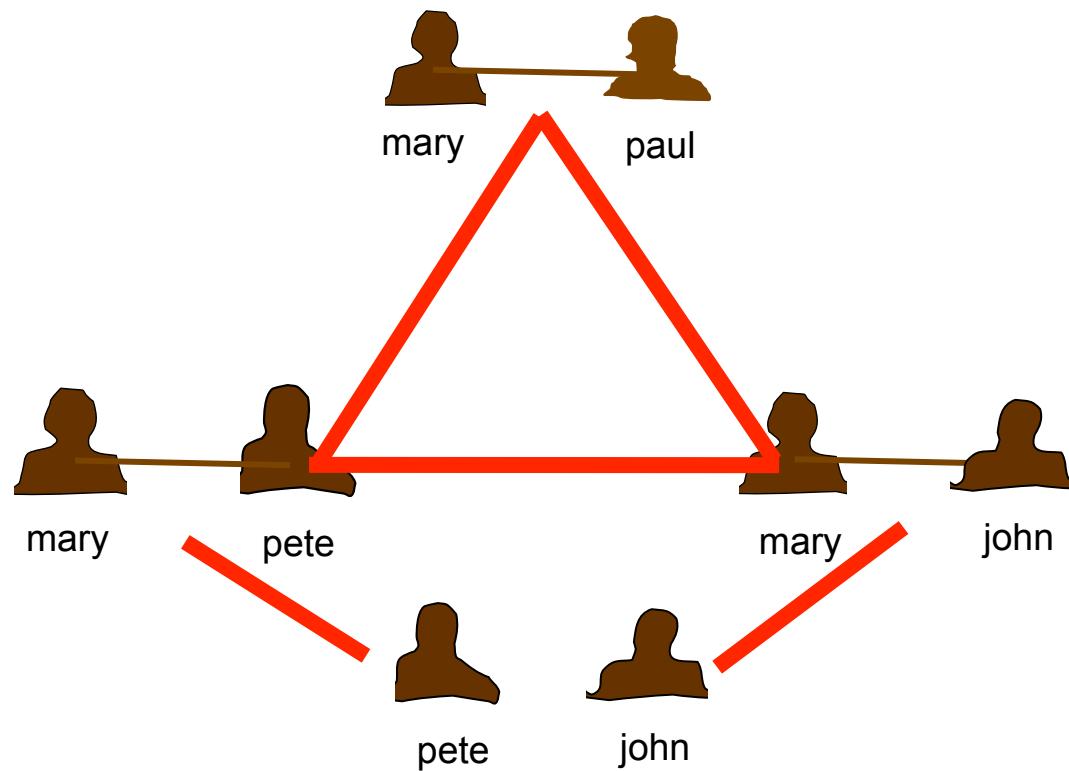
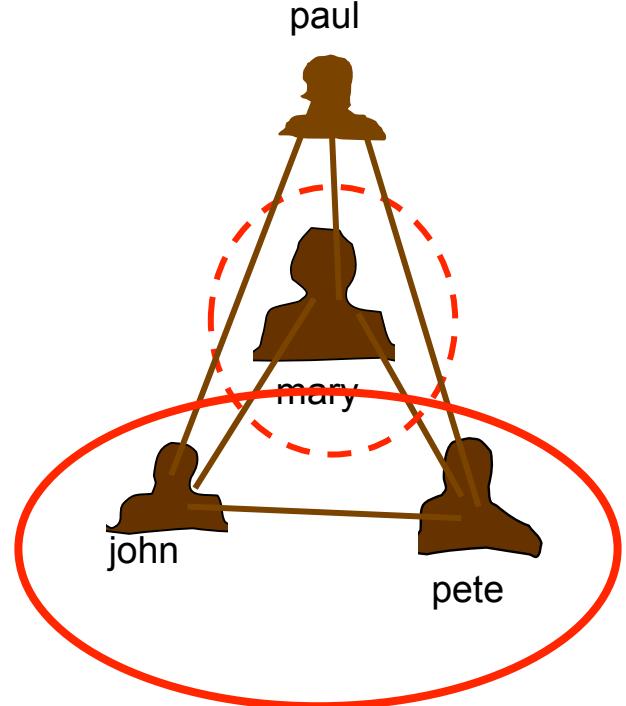
# Deriving the ERGM: From Markov graph to Dependence graph



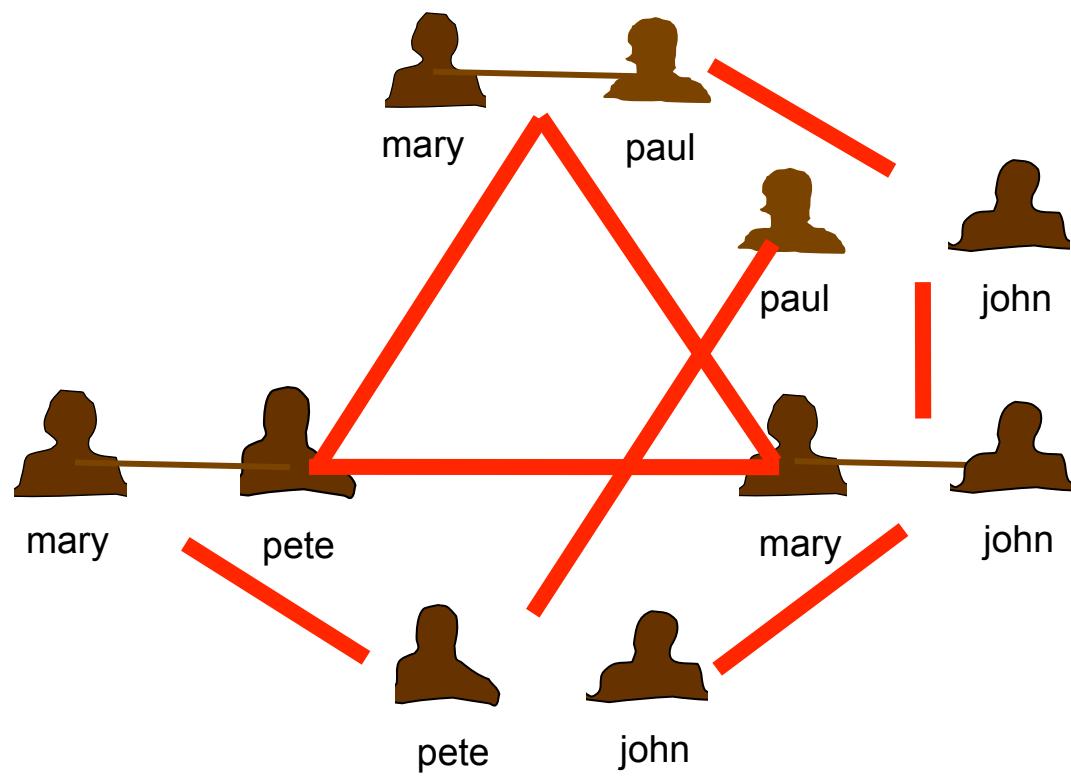
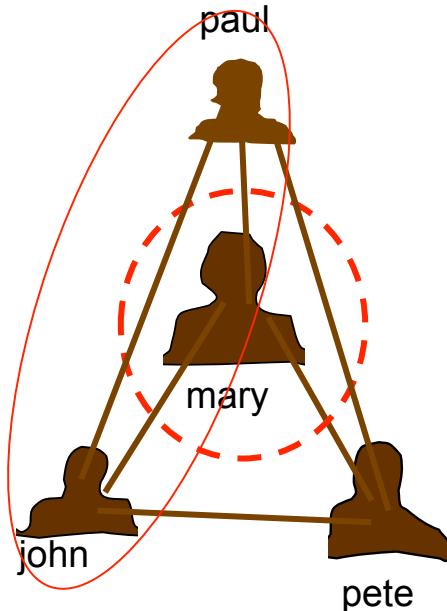
# Deriving the ERGM: From Markov graph to Dependence graph



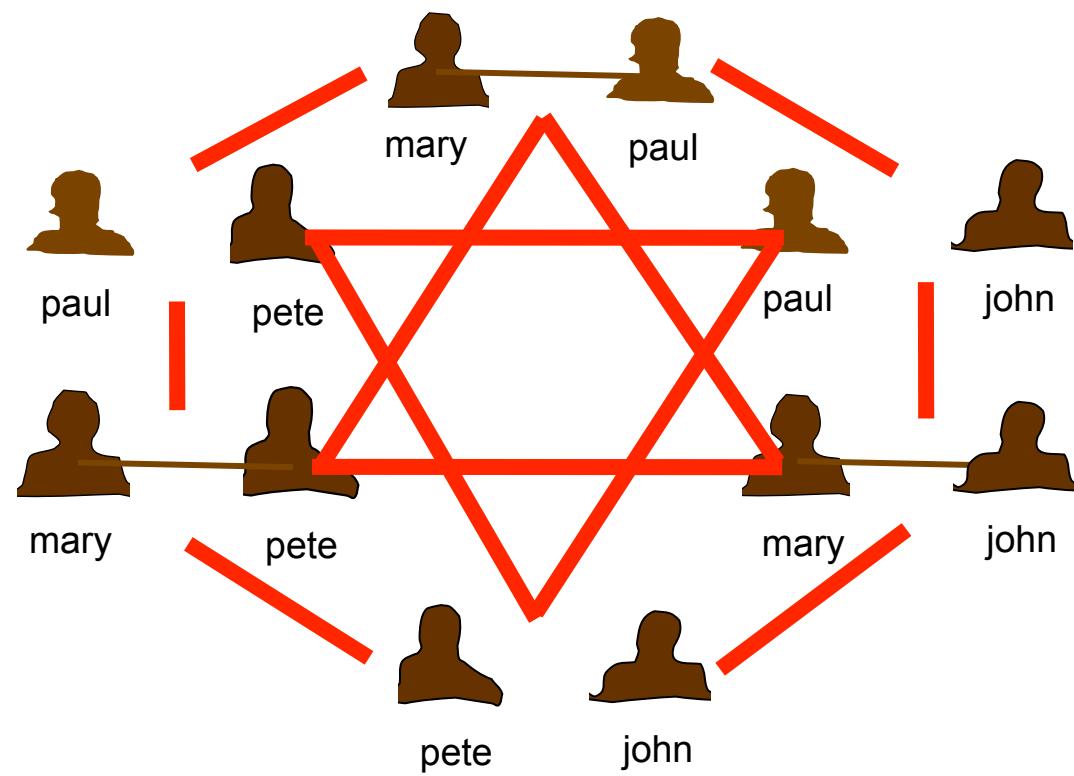
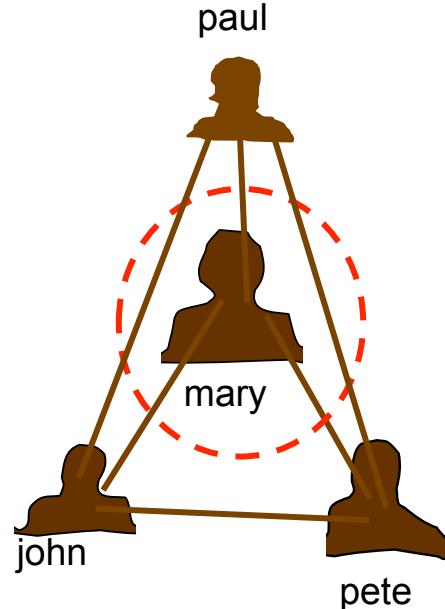
# Deriving the ERGM: From Markov graph to Dependence graph



# Deriving the ERGM: From Markov graph to Dependence graph

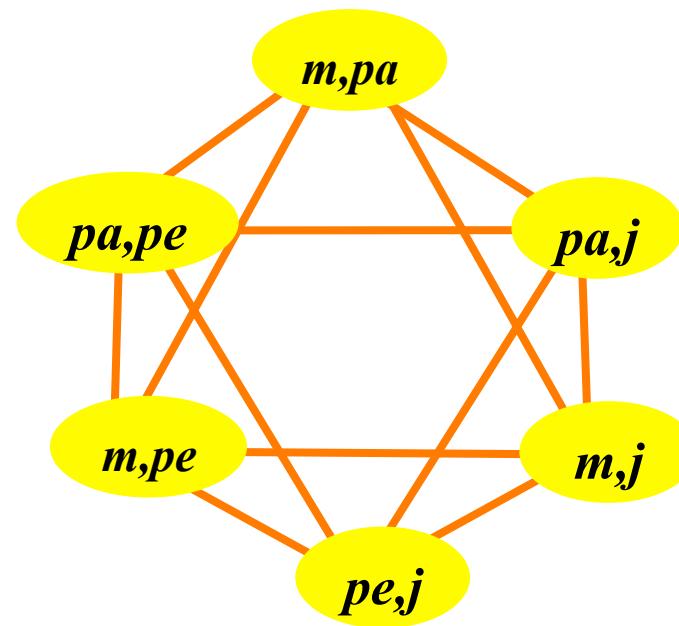
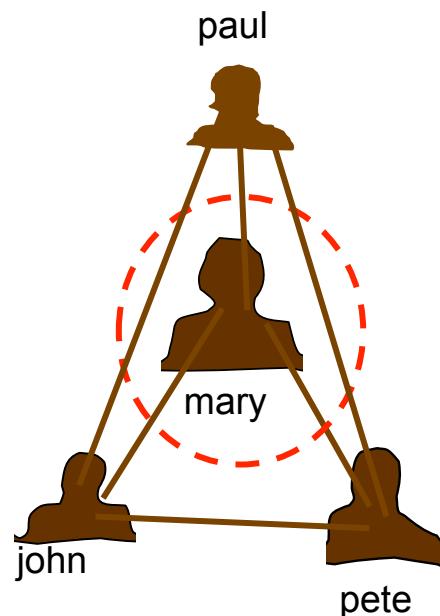


# Deriving the ERGM: From Markov graph to Dependence graph

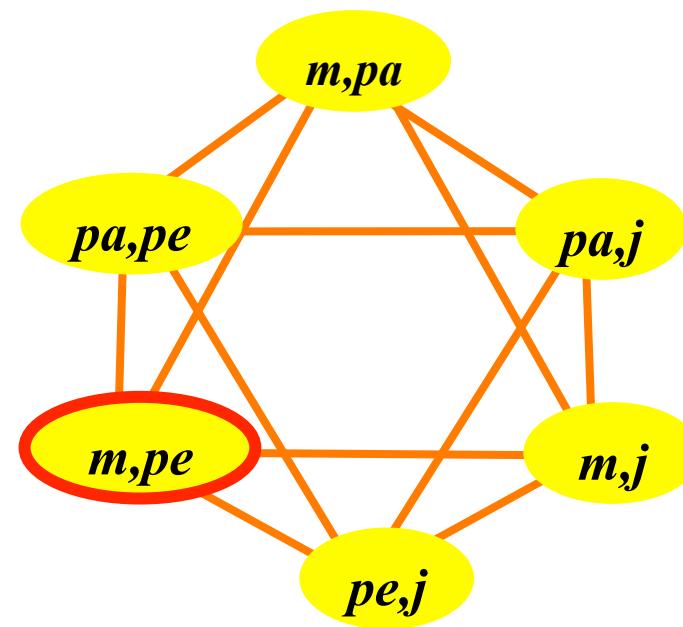
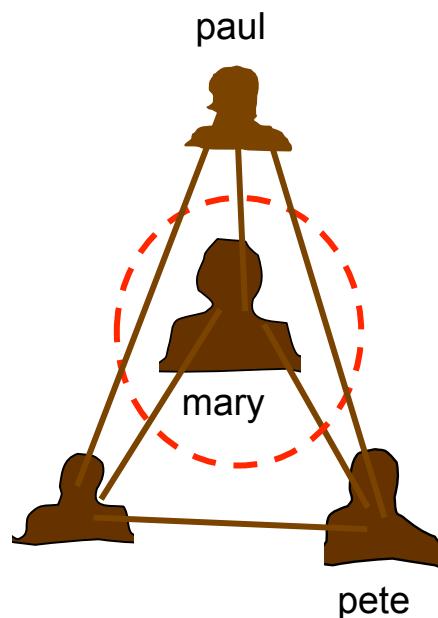


# Deriving the ERGM: From Markov graph to Dependence graph

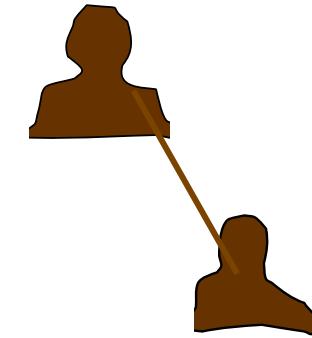
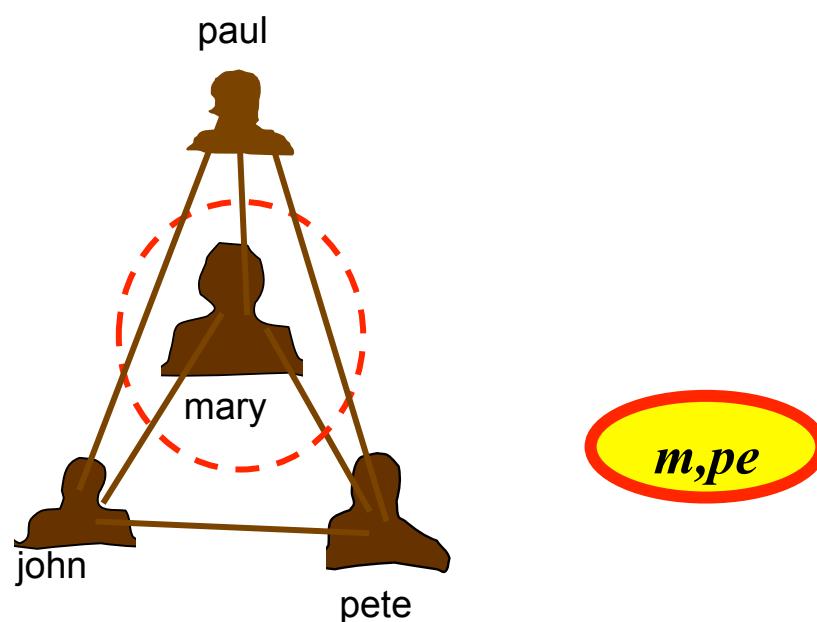
The “probability structure” of a Markov graph is described by **cliques** of the dependence graph (Hammersley-Clifford)....



# Deriving the ERGM: From Markov graph to Dependence graph

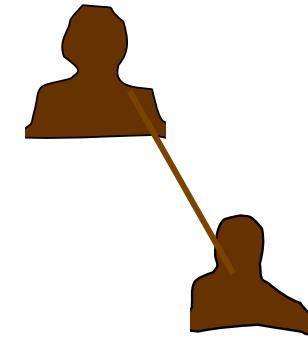
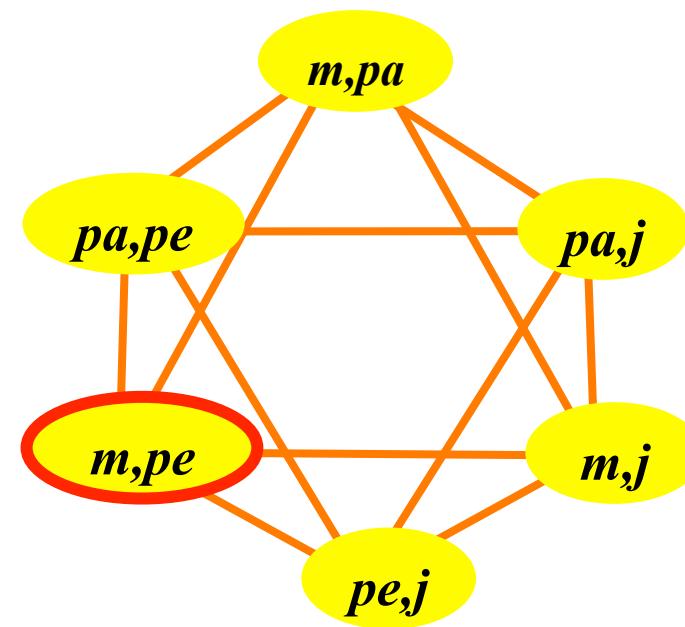
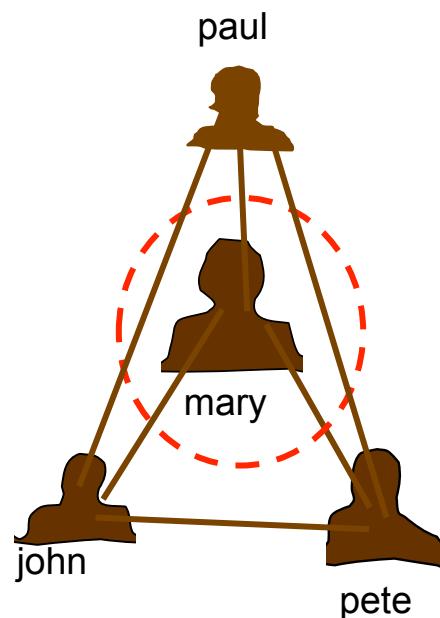


# Deriving the ERGM: From Markov graph to Dependence graph

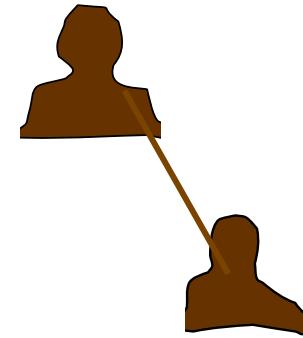
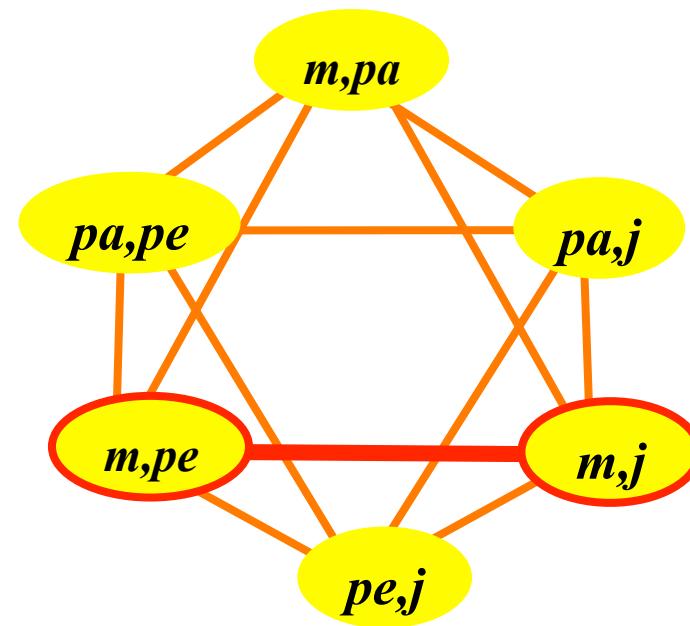
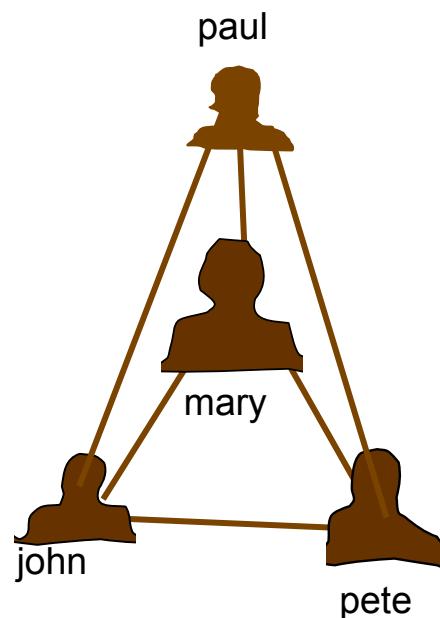


*m,pe*

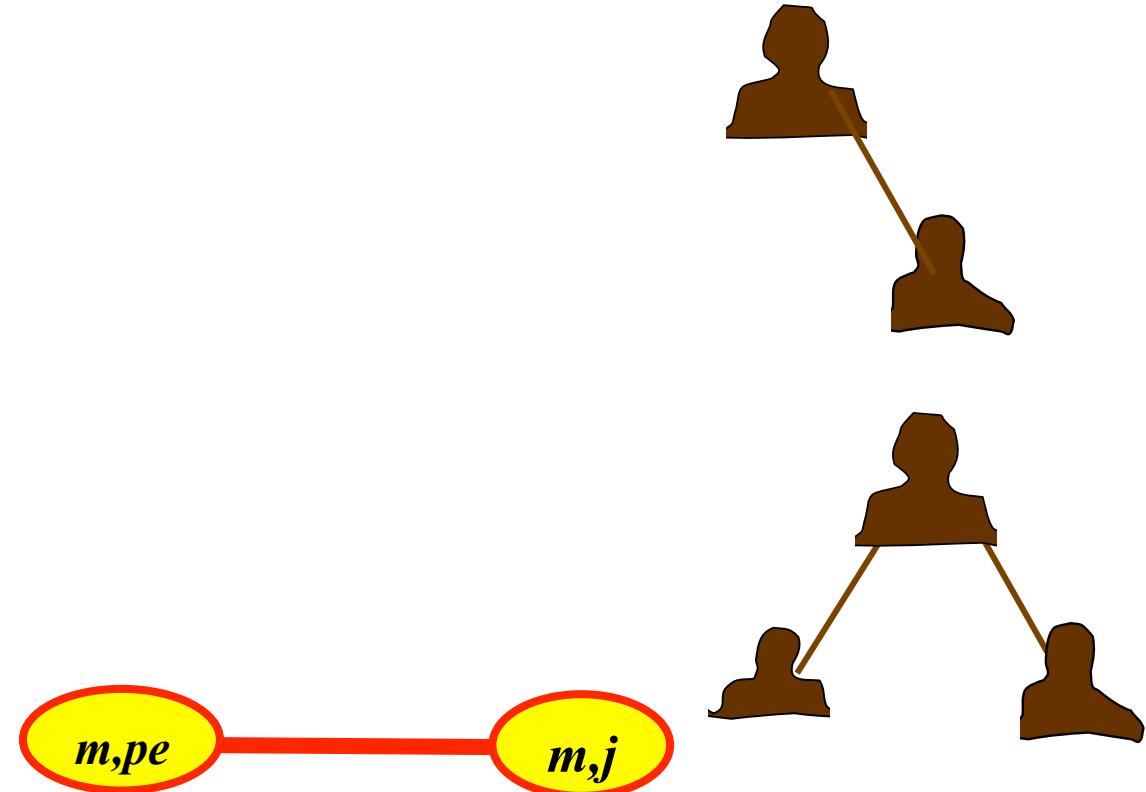
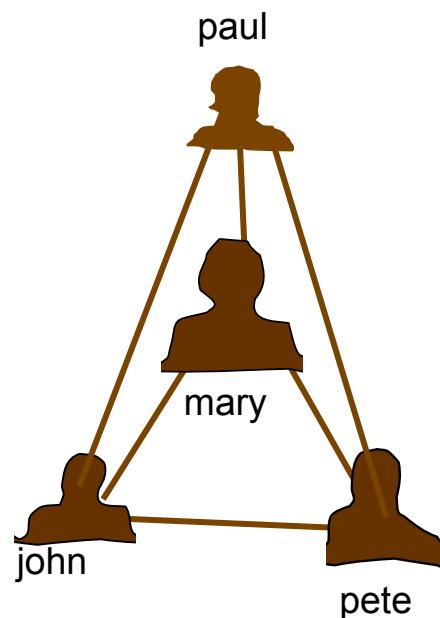
# Deriving the ERGM: From Markov graph to Dependence graph



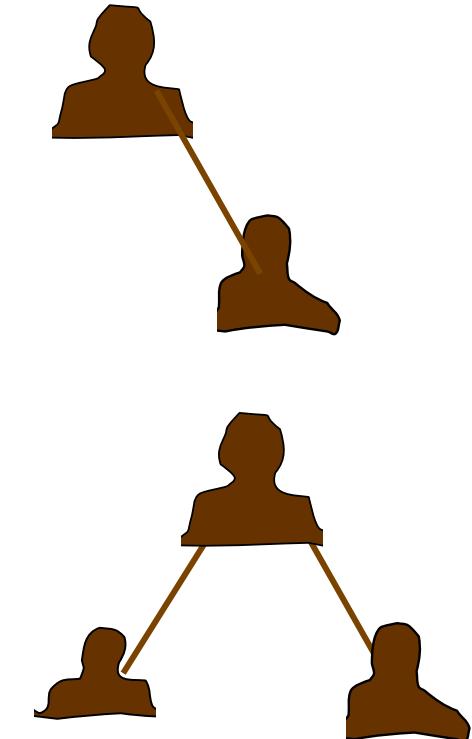
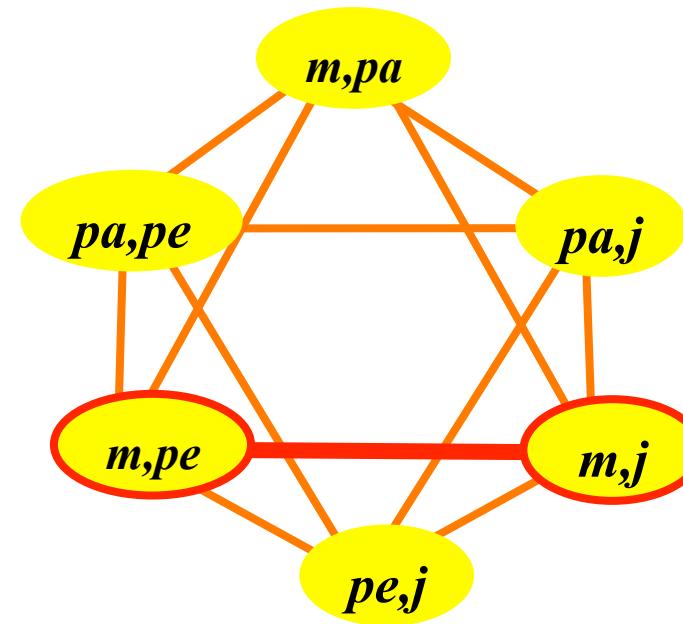
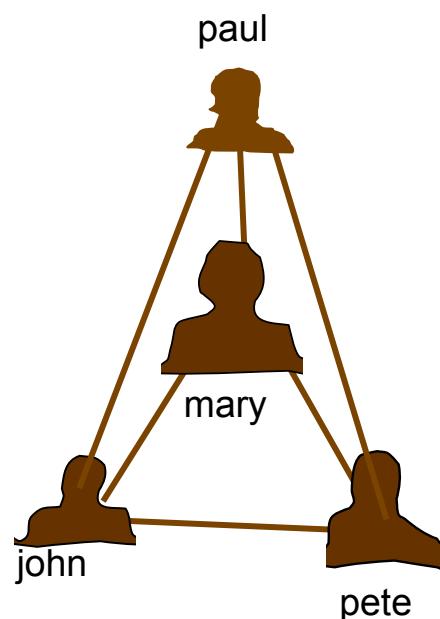
# Deriving the ERGM: From Markov graph to Dependence graph



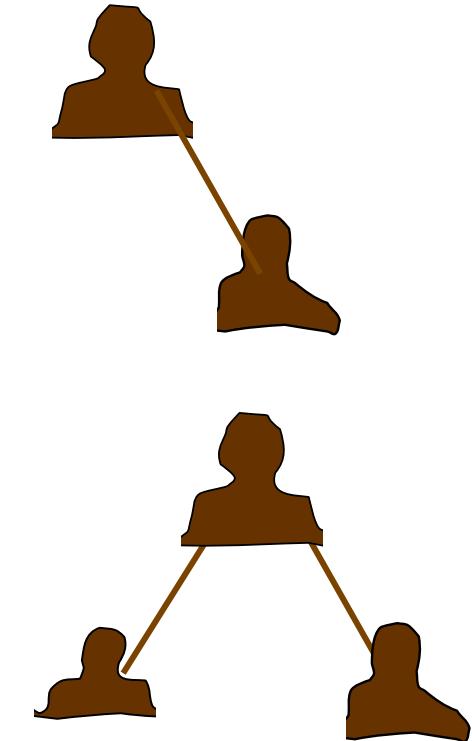
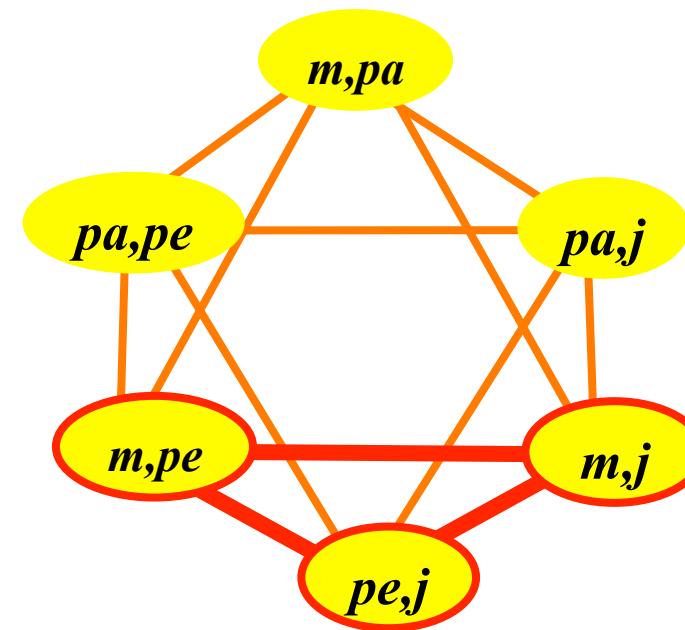
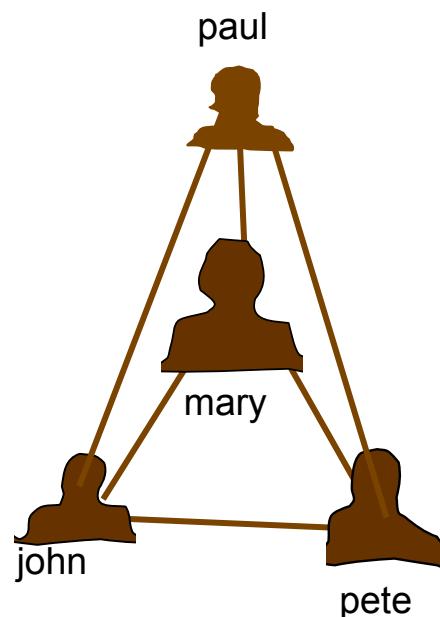
# Deriving the ERGM: From Markov graph to Dependence graph



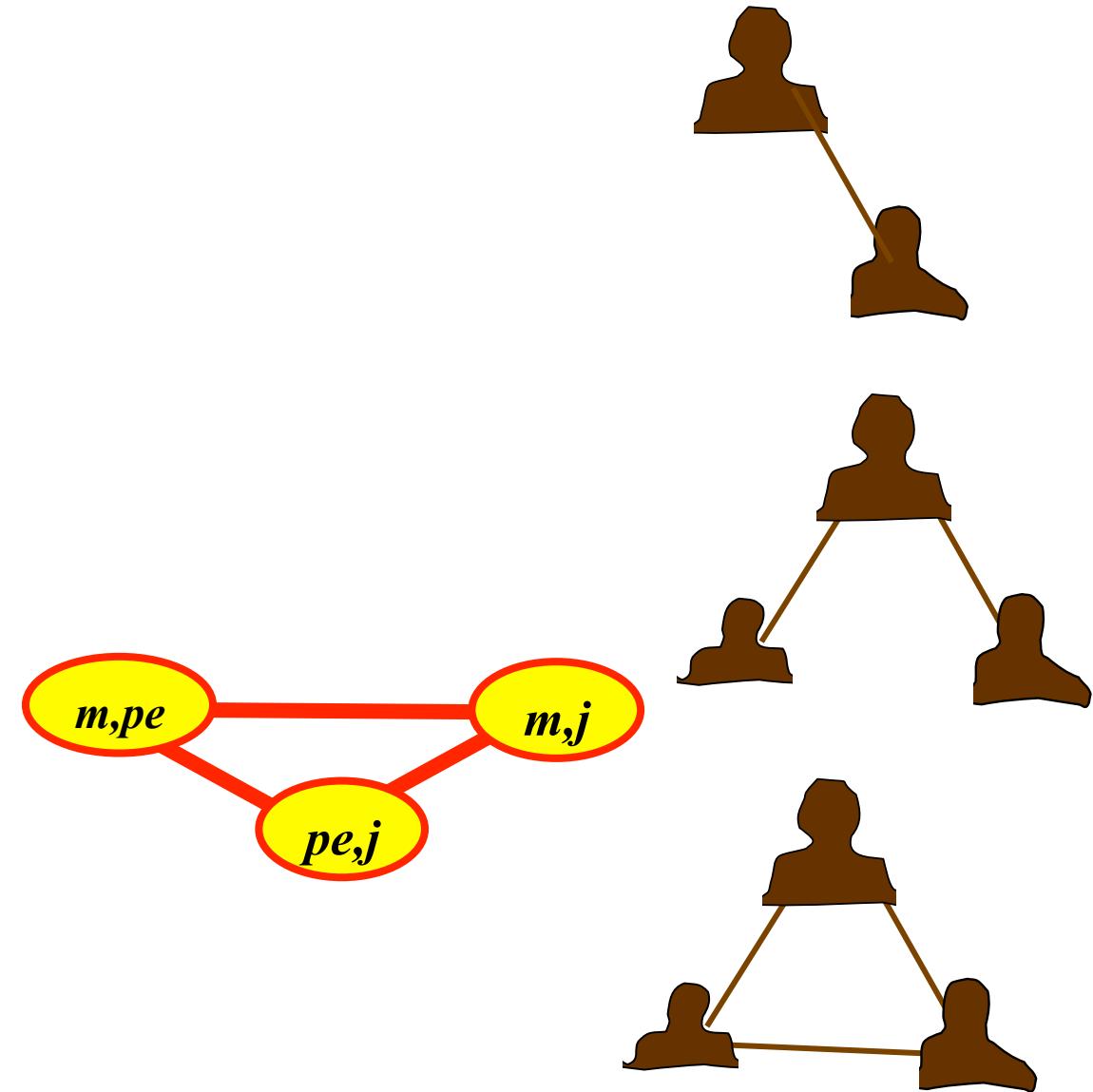
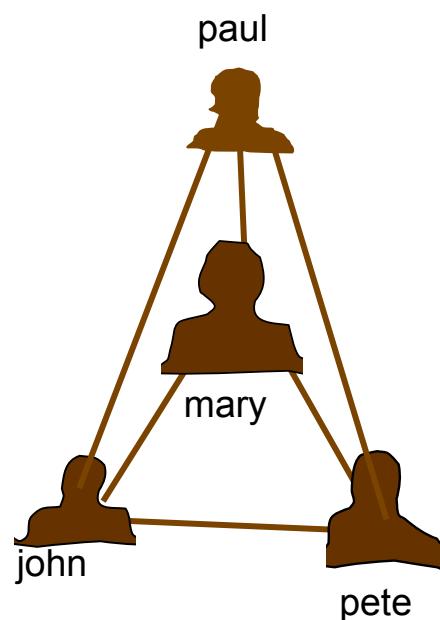
# Deriving the ERGM: From Markov graph to Dependence graph



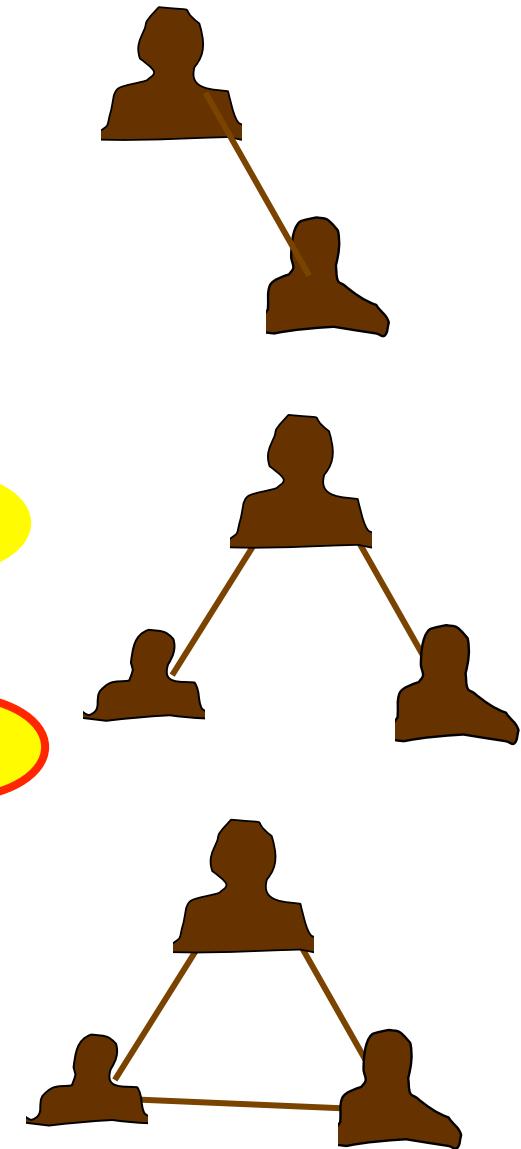
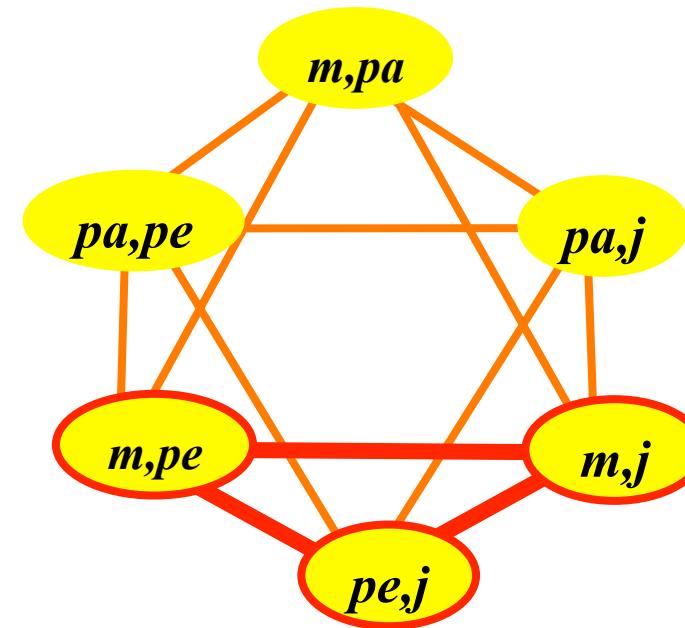
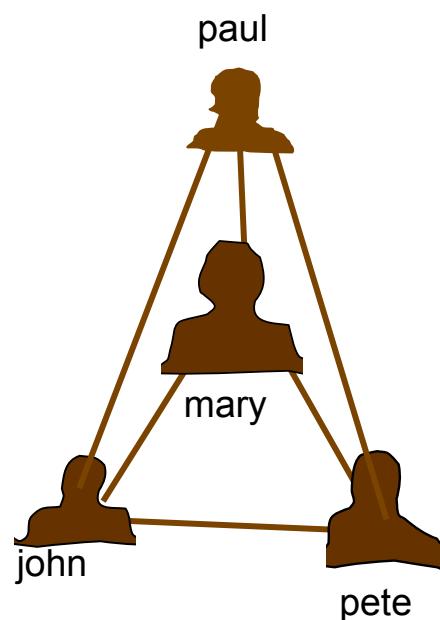
# Deriving the ERGM: From Markov graph to Dependence graph



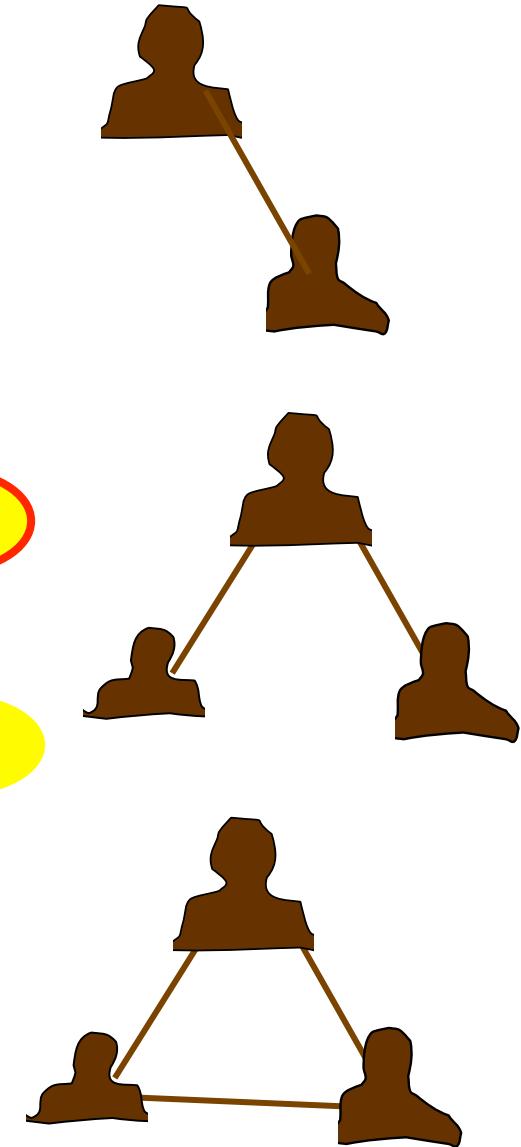
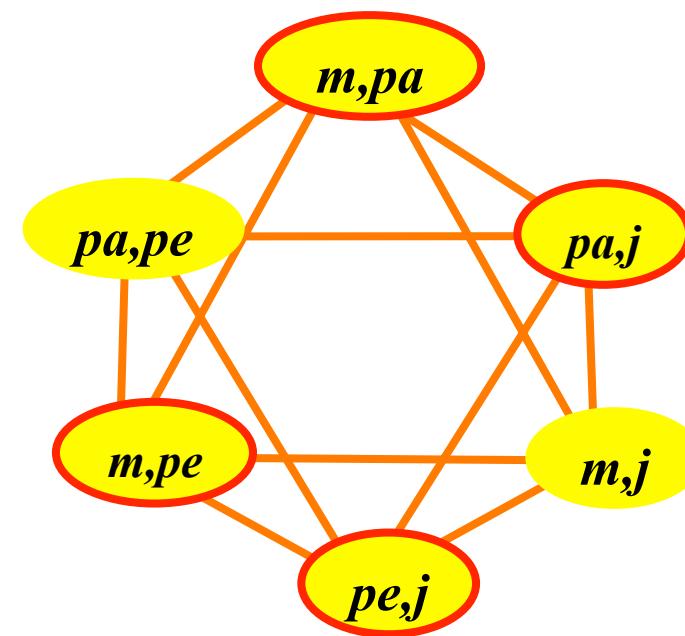
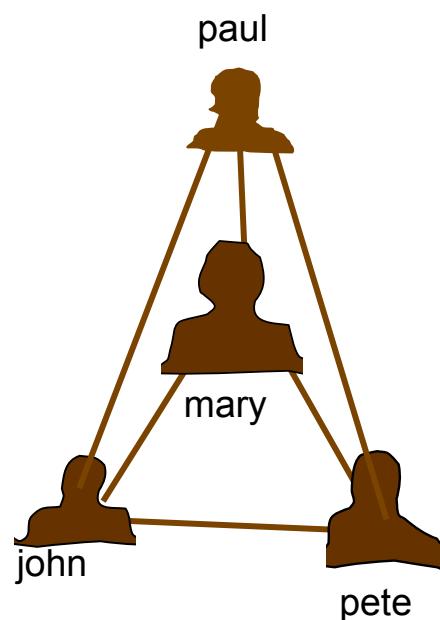
# Deriving the ERGM: From Markov graph to Dependence graph



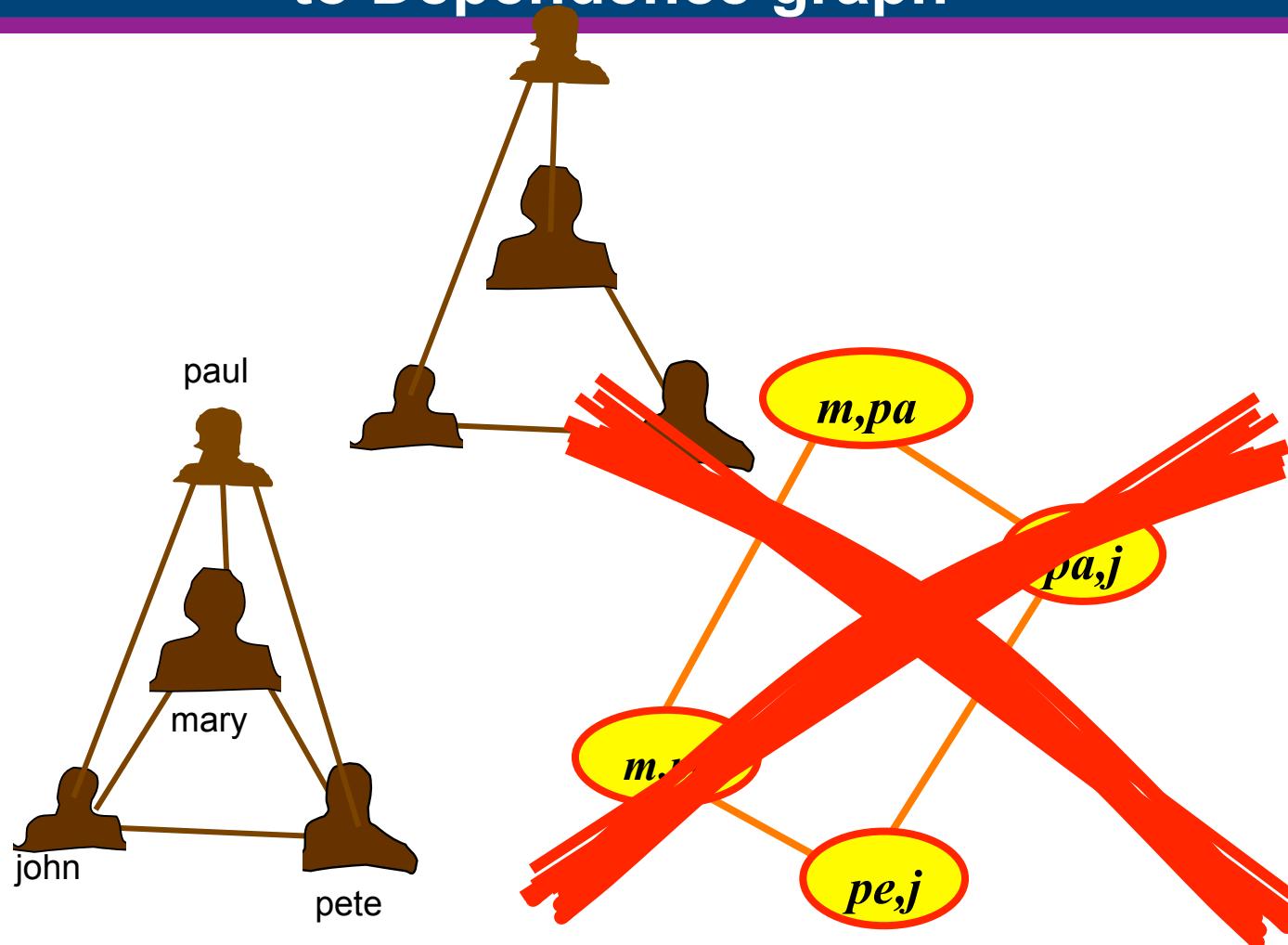
# Deriving the ERGM: From Markov graph to Dependence graph



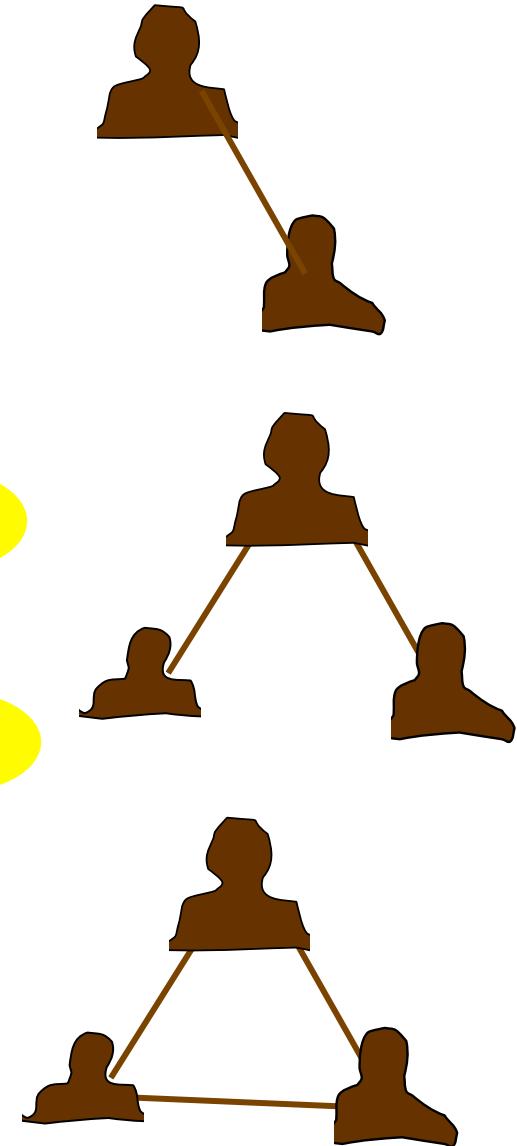
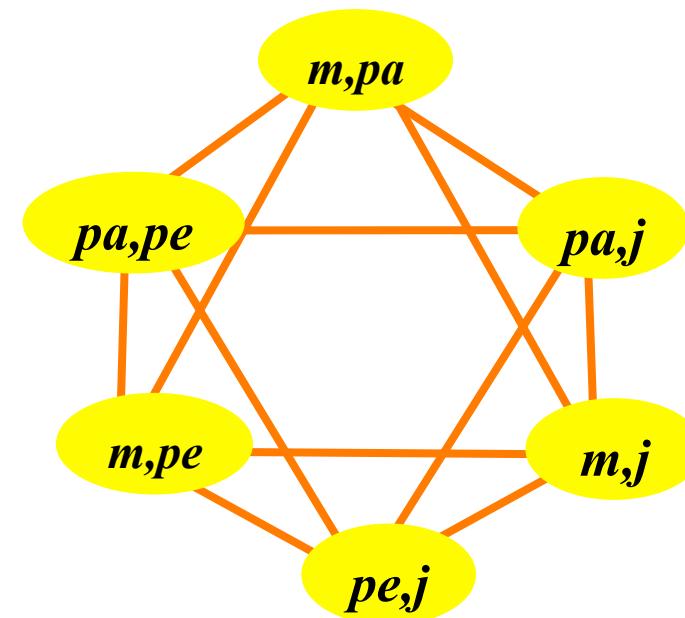
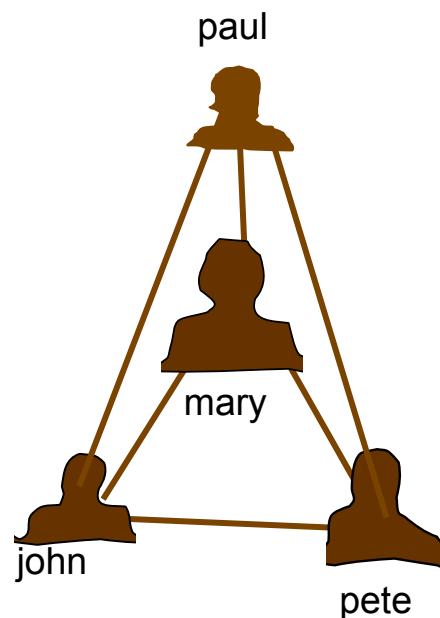
# Deriving the ERGM: From Markov graph to Dependence graph



# Deriving the ERGM: From Markov graph to Dependence graph



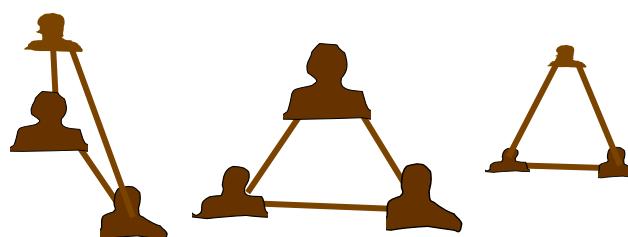
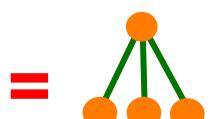
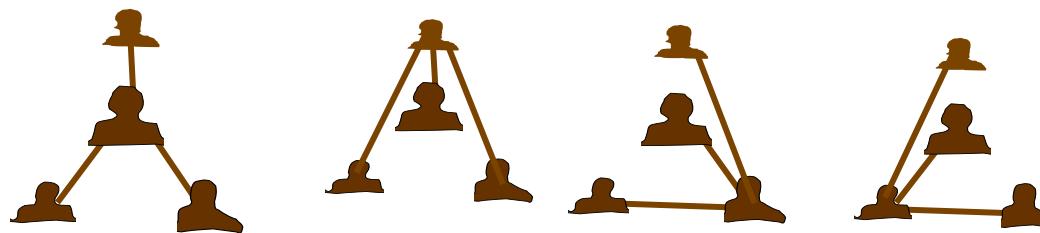
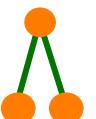
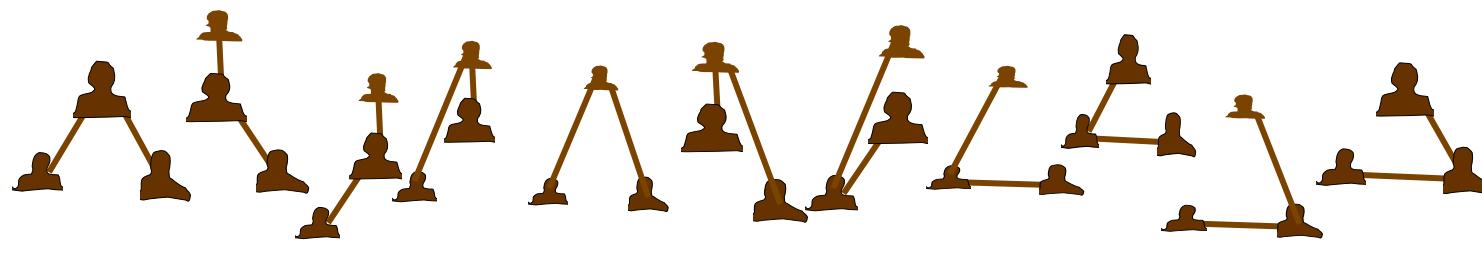
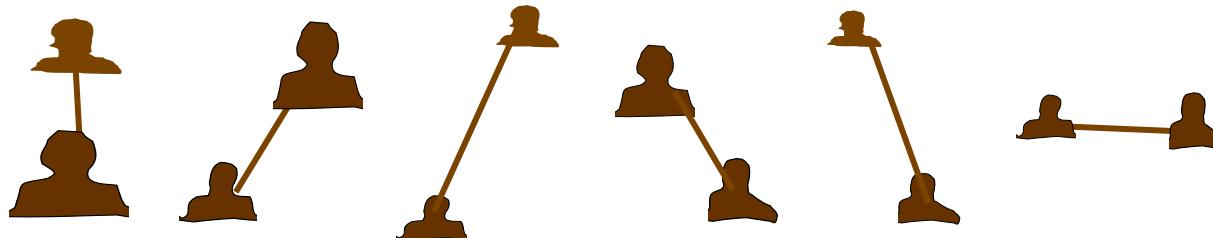
# Deriving the ERGM: From Markov graph to Dependence graph



# From Markov graph to Dependence graph – distinct subgraphs?

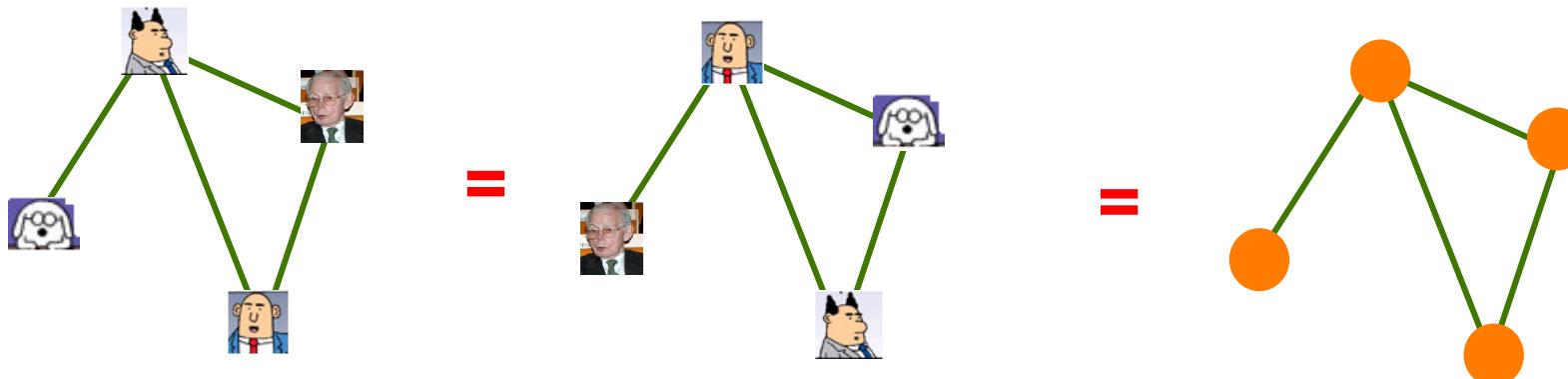


# The homogeneity assumption



# The homogeneity assumption

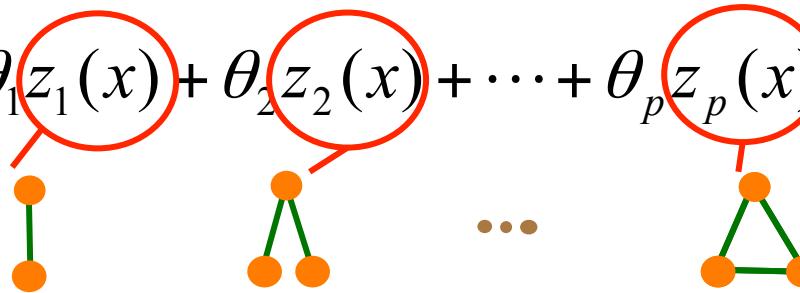
Interpretation: the probability of a graph depends only on the structure of the graph



# A log-linear model (ERGM) for ties

"Aggregated" to a joint model for **entire adjacency matrix**

$$\log \Pr(X = x) = \theta_1 z_1(x) + \theta_2 z_2(x) + \cdots + \theta_p z_p(x) + \psi(\theta)$$



Interaction terms in log-linear model of types

$$X_{ij}$$

$$X_{ij} X_{ik}$$

...

$$X_{ij} X_{ik} X_{jk}$$

# A log-linear model (ERGM) for ties

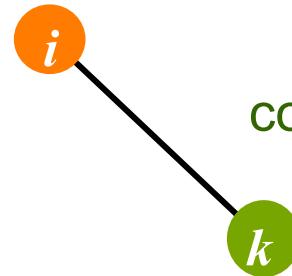
By definition of (in-) dependence

$$\Pr(X_{ij} = x_{ij}, X_{ik} = x_{ik}) \neq \Pr(X_{ij} = x_{ij}) \Pr(X_{ik} = x_{ik})$$

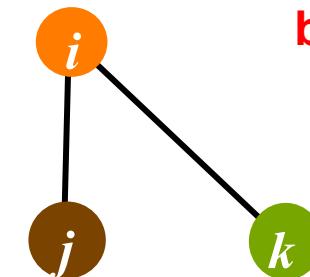
E.g.



and



co-occurring



**More than is explained  
by margins**

$$X_{ij}$$

**Main effects**

$$X_{ik}$$

$$X_{ij} X_{ik}$$

**interaction term**

## Part 7

### Summary of fitting routine

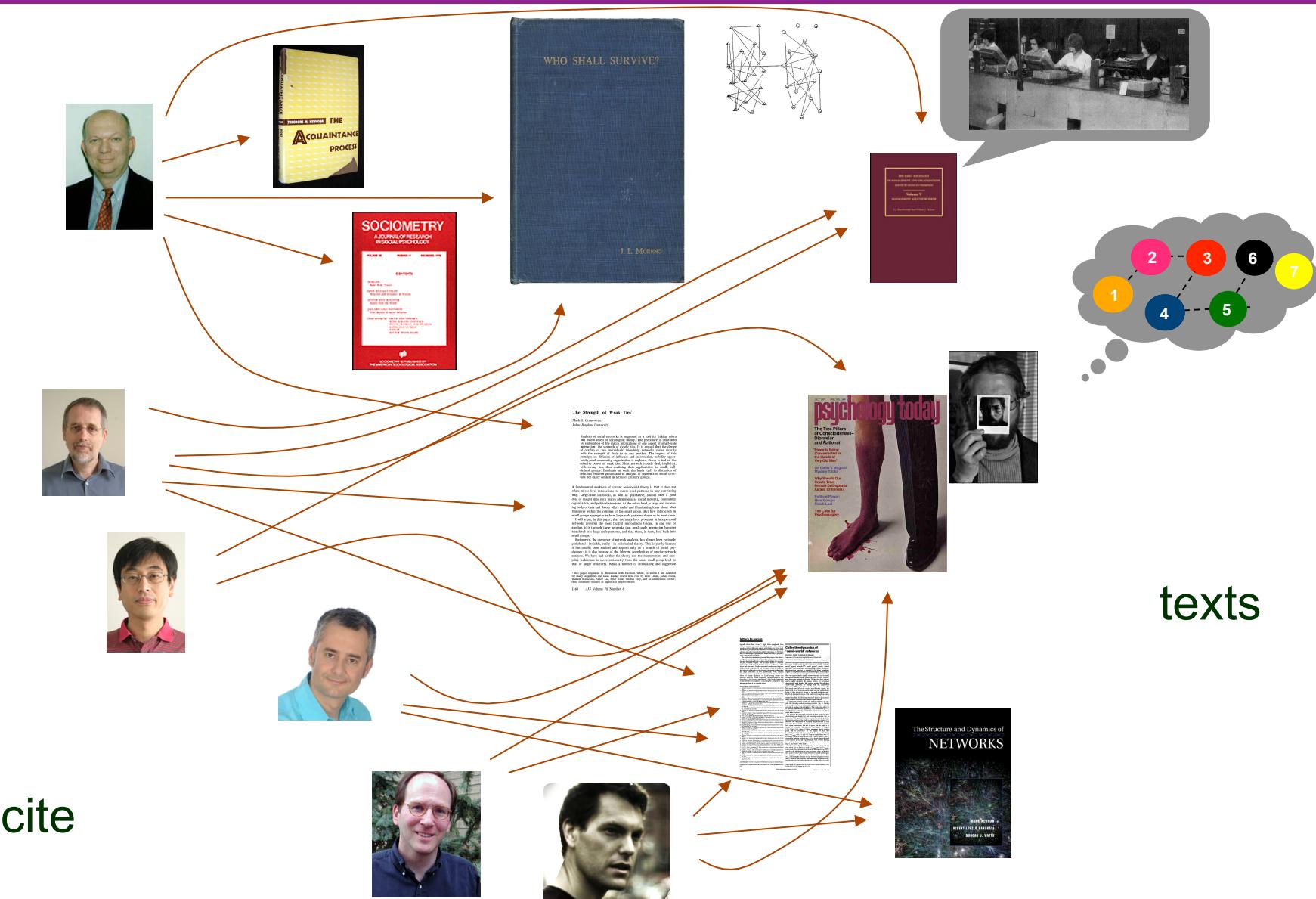
## The steps of fitting an ERGM

- fit base-line model
- check convergence
- rerun if model not converged
- include more parameters? GOF
- candidate models

# Part 8

## Bipartite data

# Bi-partite networks: cocitation (Small 1973)



# Bi-partite networks: cooffending (. Sarnecki, 2001)

offenders



offences

participating

# Bi-partite networks

people



participating in

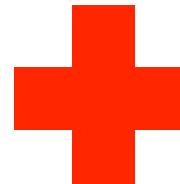


Social events  
(Breiger, 1974)

people

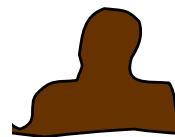


belonging to



voluntary organisations  
(Bonachich, 1978)

directors



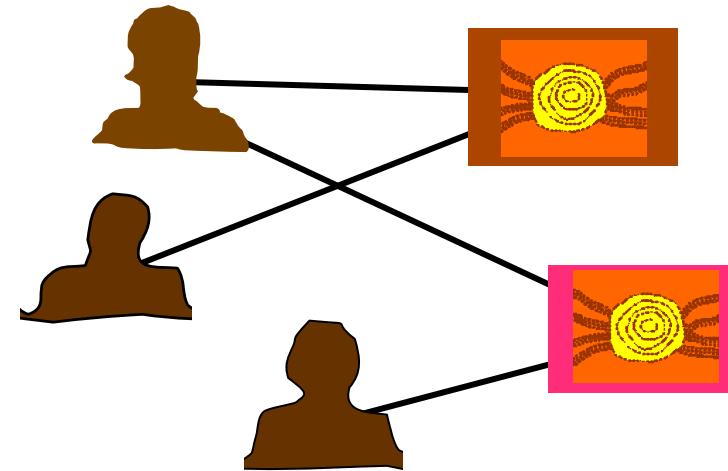
sitting on



corporate boards  
(eg. Mizruchi, 1982)

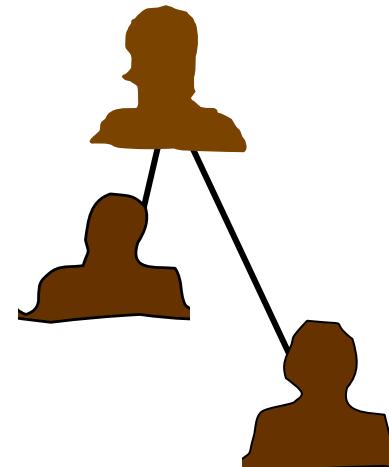
# One-mode projection

Two-mode

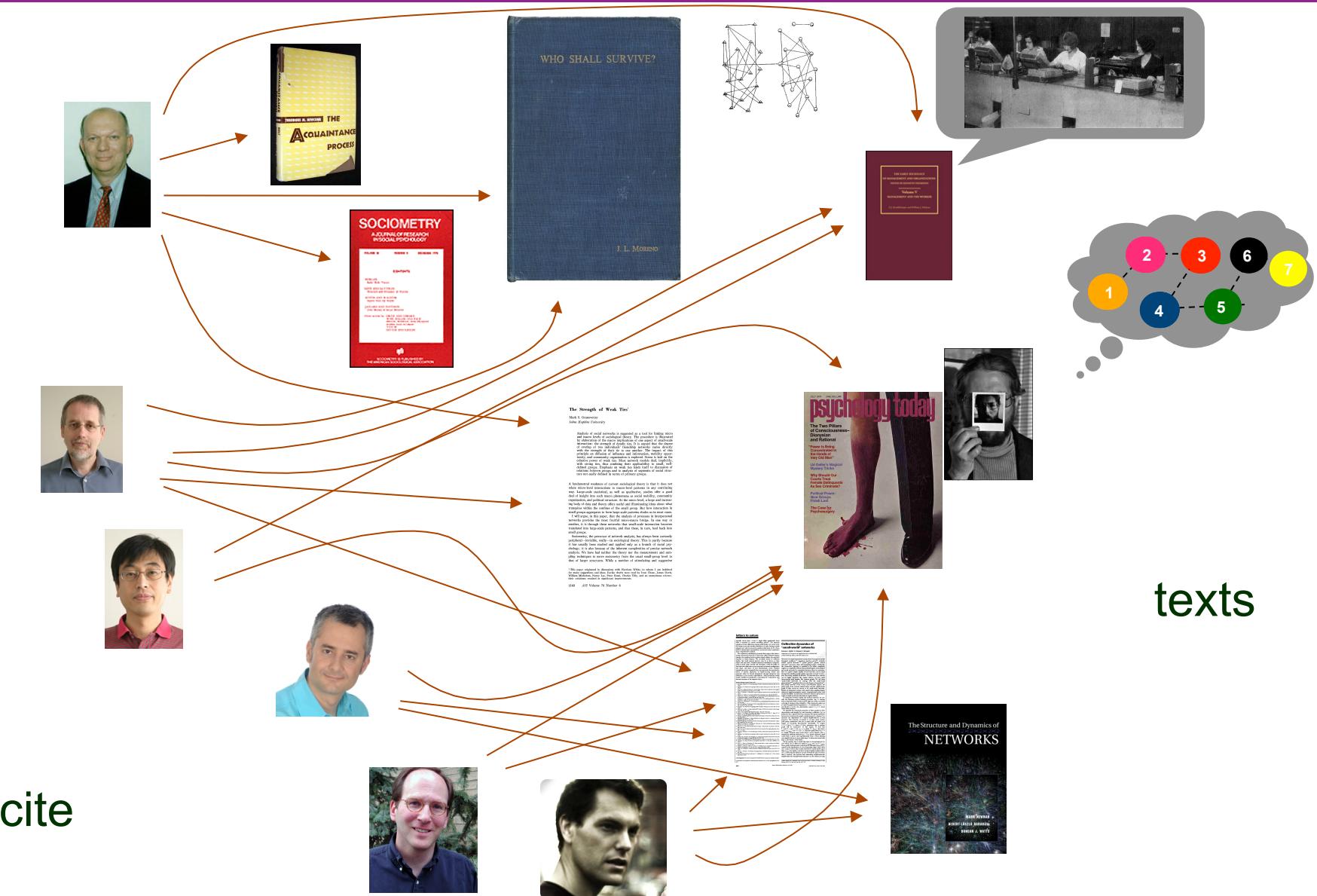


**Tie:** If two directors share a board

one-mode

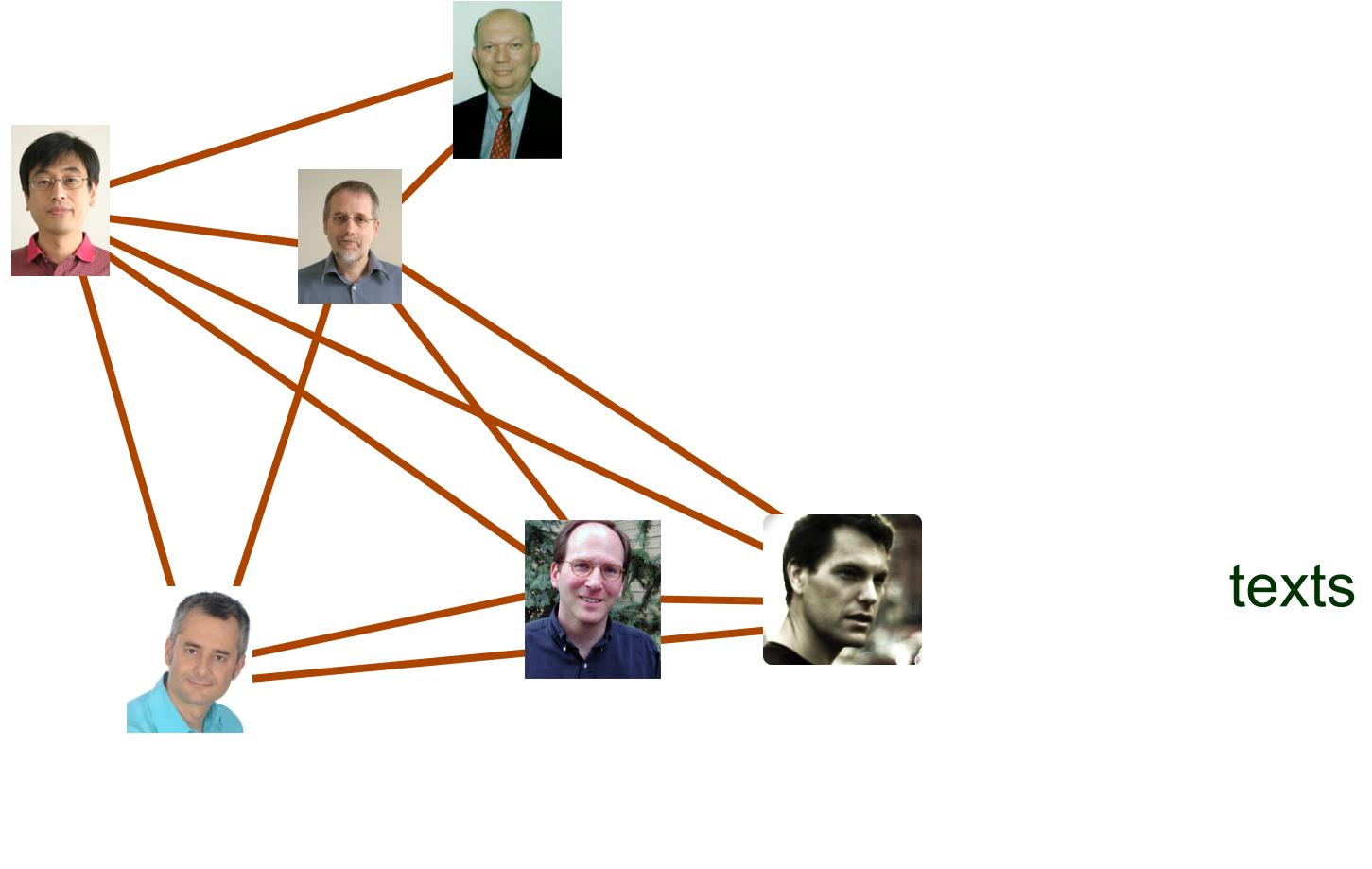


# Bi-partite networks



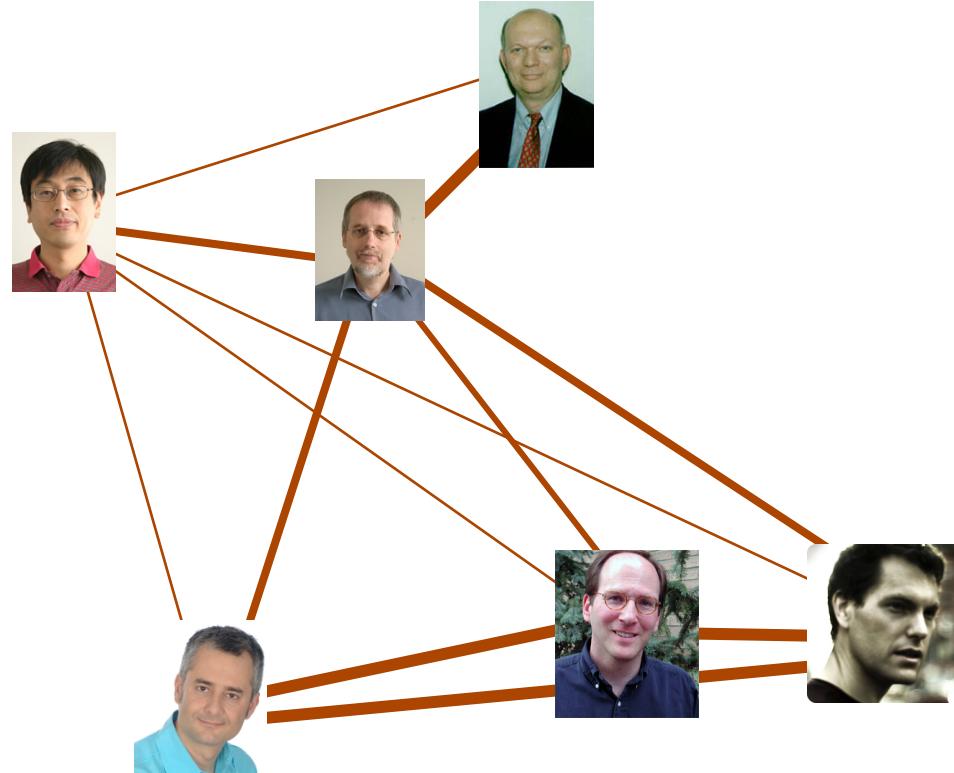
# Bi-partite networks

Researchers



# Bi-partite networks

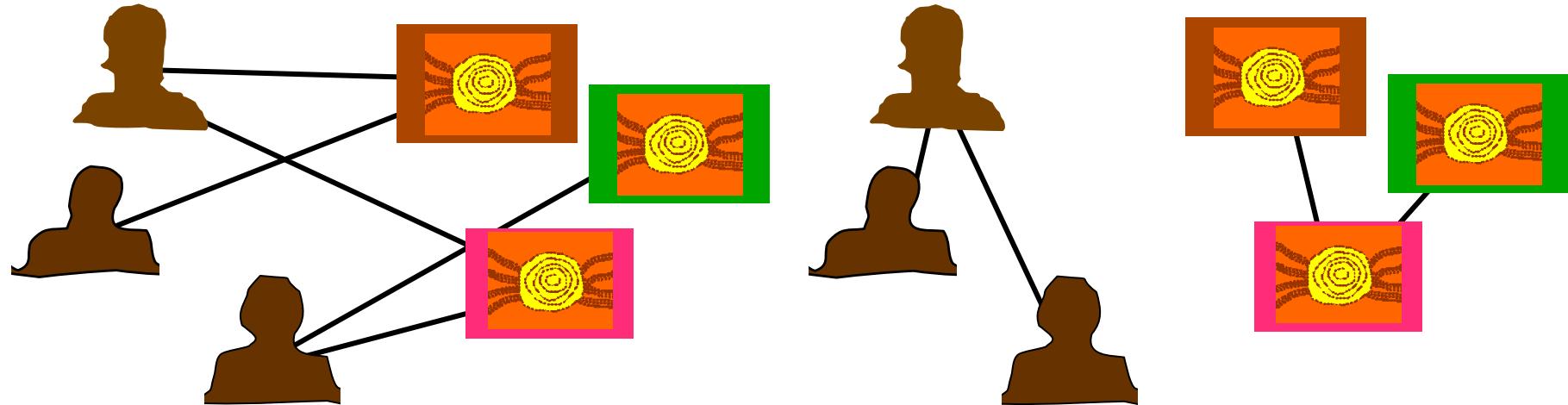
## Researchers



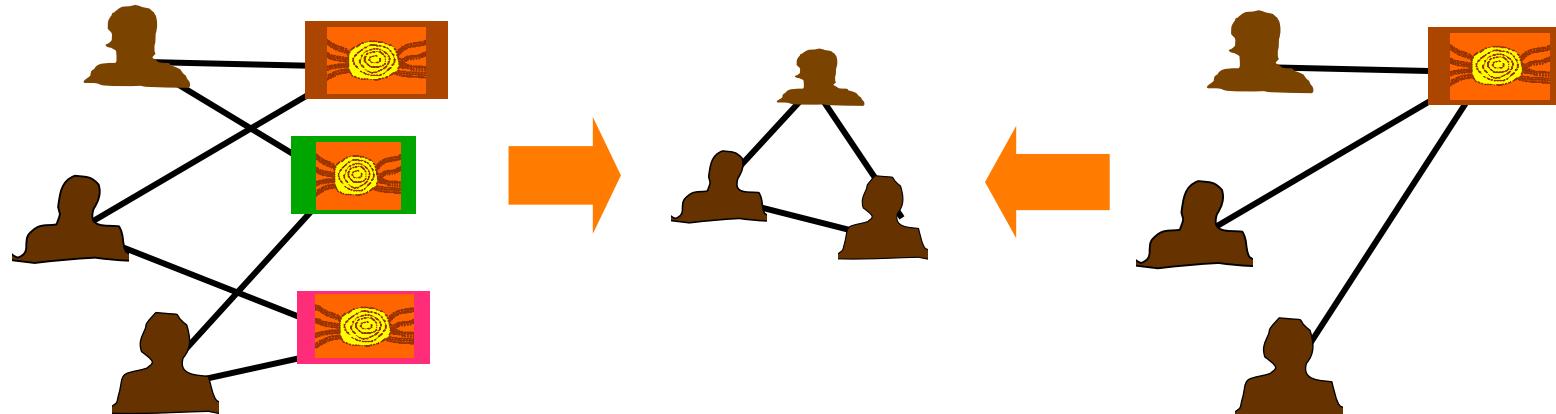
cite

texts

# One-mode projection



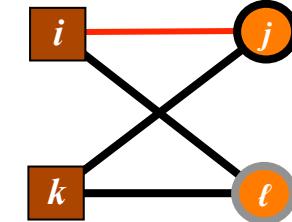
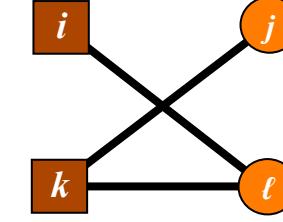
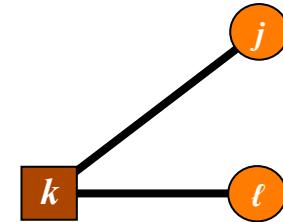
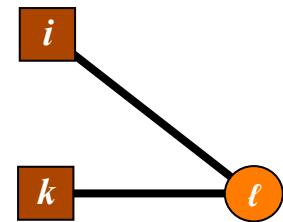
**What mode given priority?** (Duality of social actors and social groups; e.g. Breiger 1974; Breiger and Pattison, 1986)  
**Loss of information**



# ERGM for bipartite networks

The model is the same as for one-mode networks (Wang et al., 2007)

$$\log \Pr(X = x) = \theta_1 z_1(x) + \theta_2 z_2(x) + \cdots + \theta_p z_p(x) + \psi(\theta)$$



Edges

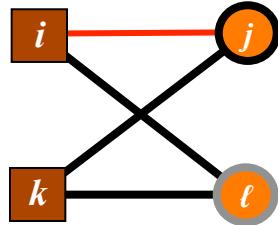
"people"  
2-stars

"affiliation"  
2-stars

3-paths

4-cycles

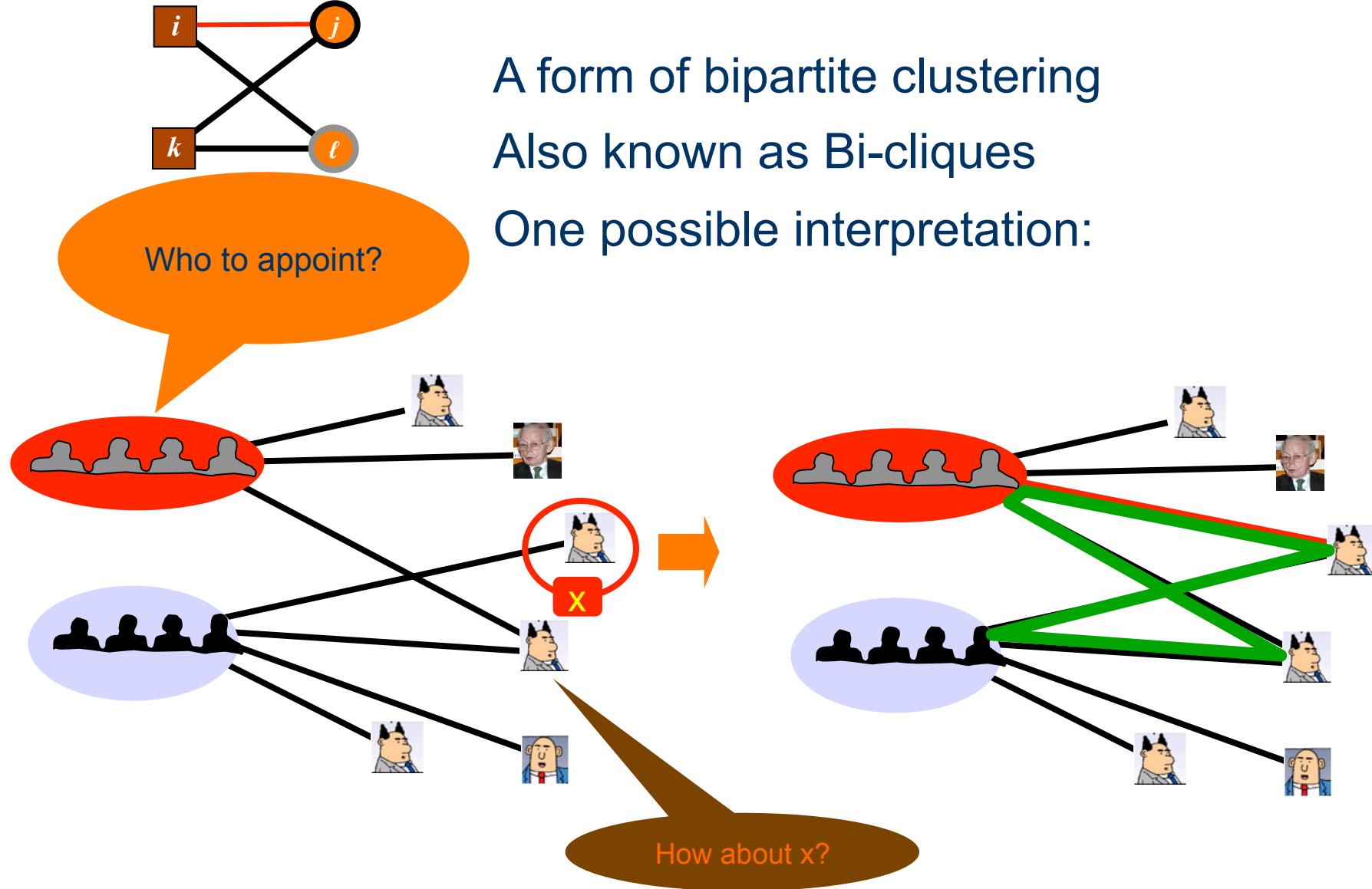
# ERGM for bipartite networks



4-cycles

A form of bipartite clustering  
Also known as Bi-cliques  
One possible interpretation:

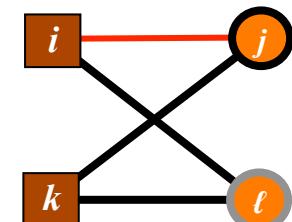
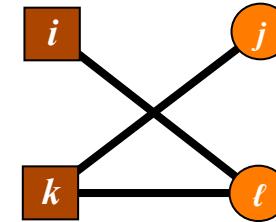
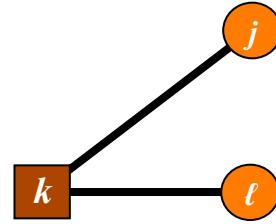
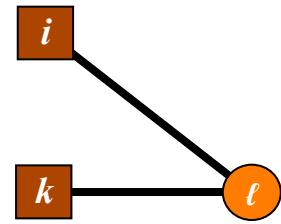
# ERGM for bipartite networks



# ERGM for bipartite networks

Fitting the model in (B)Pnet straightforward extension

These statistics are **all Markov**:



Edges

"people"  
2-stars

"affiliation"  
2-stars

3-paths

4-cycles

<BPNet>

# Part 9

## Missing data

## Effects of missingness

### Perils

Some investigations on the effects on indices of structural properties (Kossinets, 2006; Costenbader & Valente, 2003; Huisman, 2007)

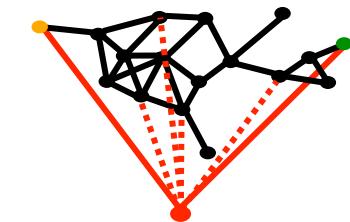
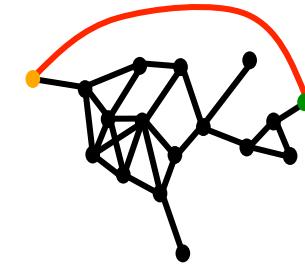
Problems with the “boundary specification issue”

### Few remedies

Deterministically “complement” data (Stork & Richards, 1992)

Stochastically Impute missing data (Huisman, 2007)

Ad-hoc “likelihood” (score) for missing ties (Liben-Nowell and Kleinberg, 2007)



# Model assisted treatment of missing network data

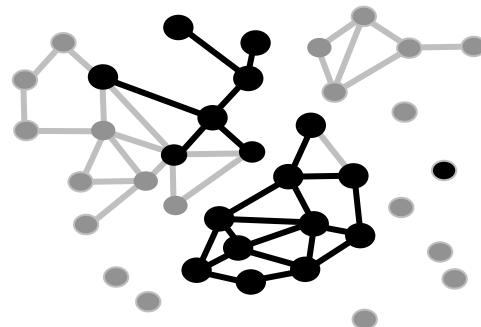
If you don't have a model for what you **have observed**

How are you going to be able to say something about what you **have not observed using** what you **have observed**

# Model assisted treatment of missing network data

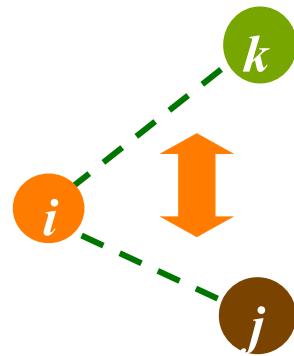
- Importance sampling (**Handcock & Gile 2010; Koskinen, Robins & Pattison, 2010**)
- Stochastic approximation and the missing data principle (**Orchard & Woodbury, 1972**)  
(Koskinen & Snijders, forthcoming)
- Bayesian data augmentation (**Koskinen, Robins & Pattison, 2010**)

# Marginalisation (Snijders, 2010; Koskinen et al, 2010)



Subgraph of ERGM not ERGM

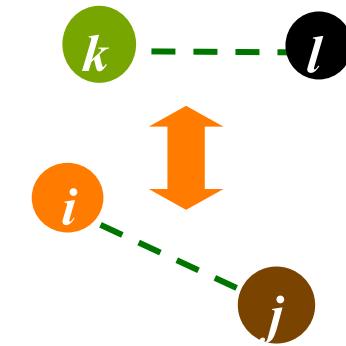
Dependence in ERGM



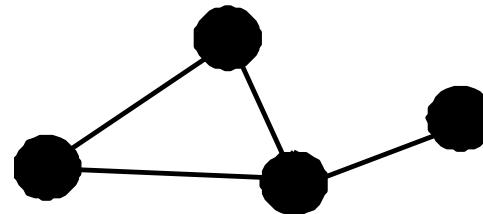
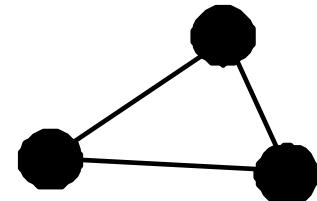
But if



We may also have dependence

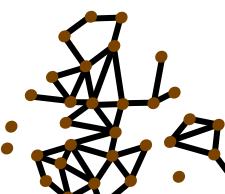
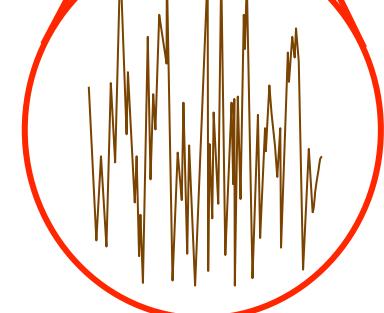
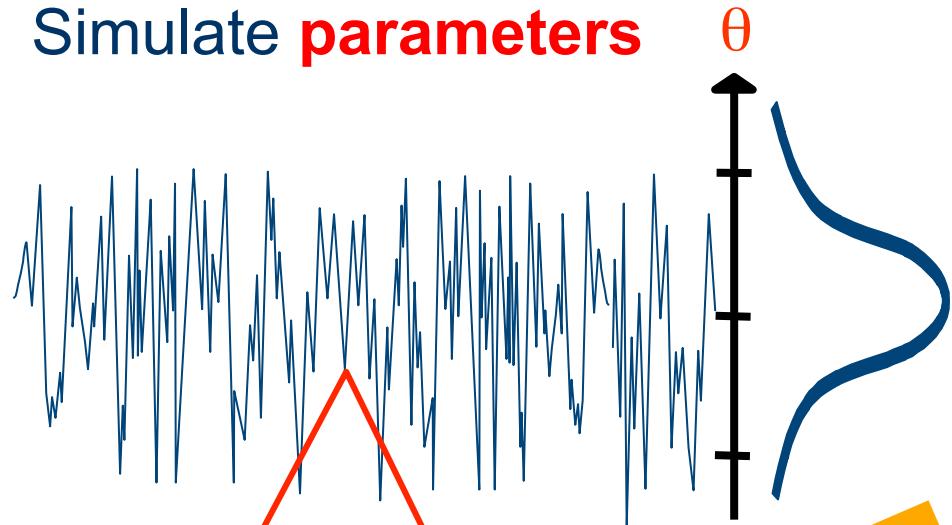


We should include counts of:



# Bayesian Data Augmentation

Simulate **parameters**



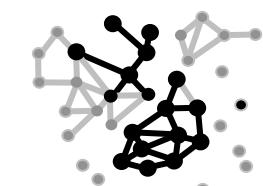
$\theta$

With missing data:

$\theta$

.

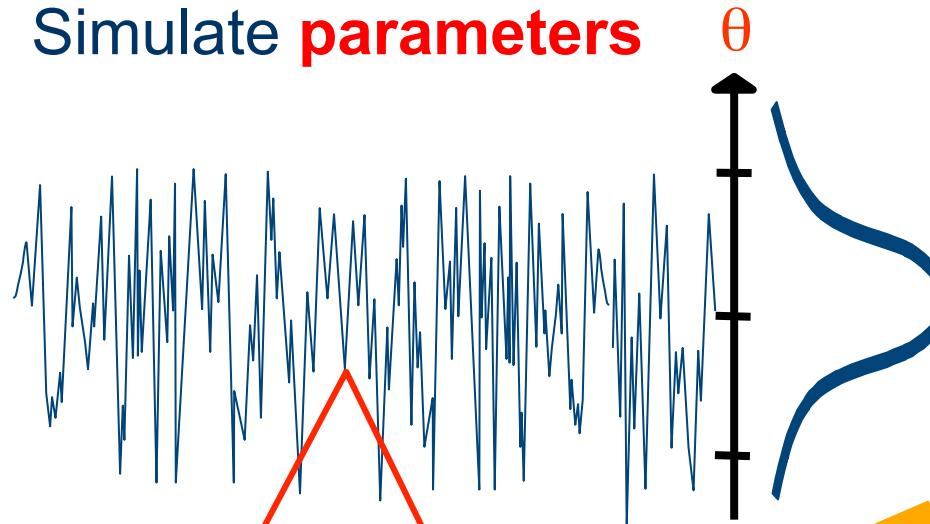
missing



In each iteration  
simulate **graphs**

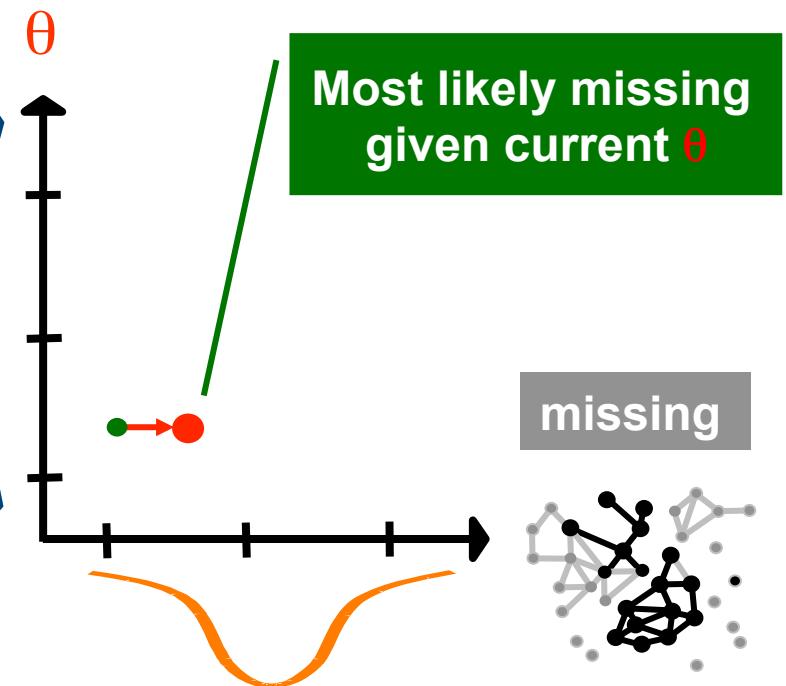
# Bayesian Data Augmentation

Simulate **parameters**



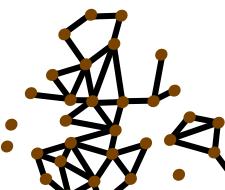
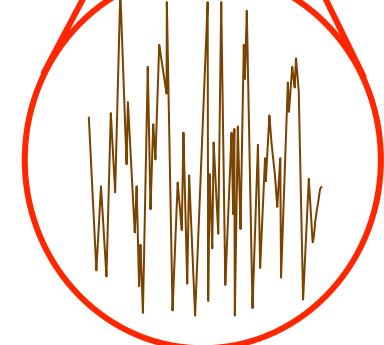
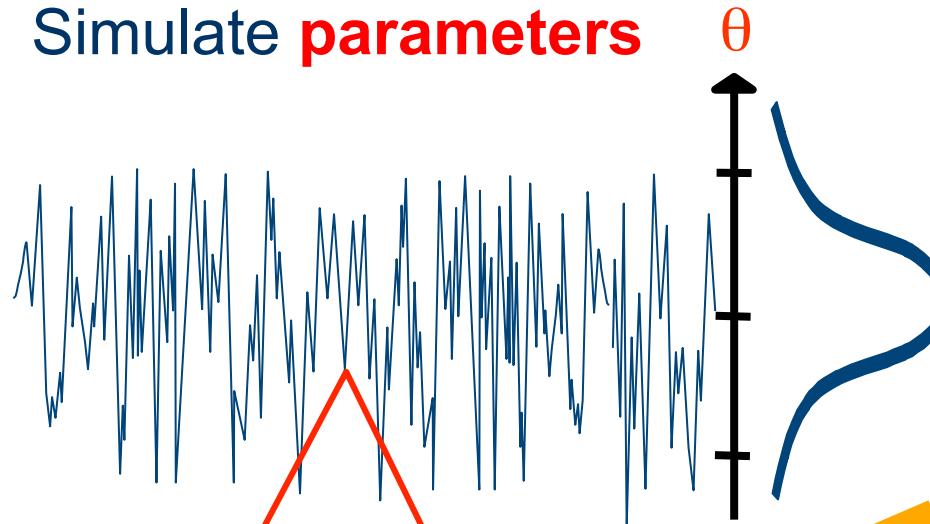
In each iteration  
simulate **graphs**

With missing data:



# Bayesian Data Augmentation

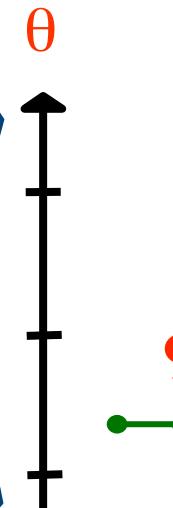
Simulate **parameters**



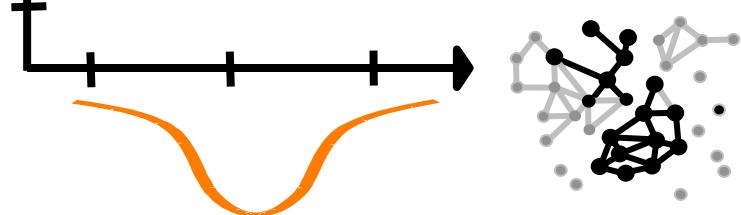
In each iteration  
simulate **graphs**

With missing data:

Most likely  $\theta$  given  
current missing

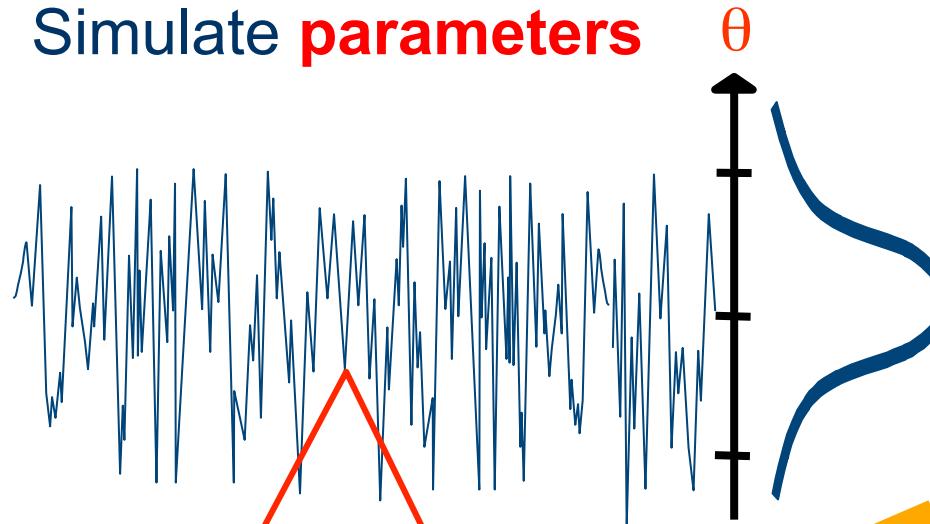


missing



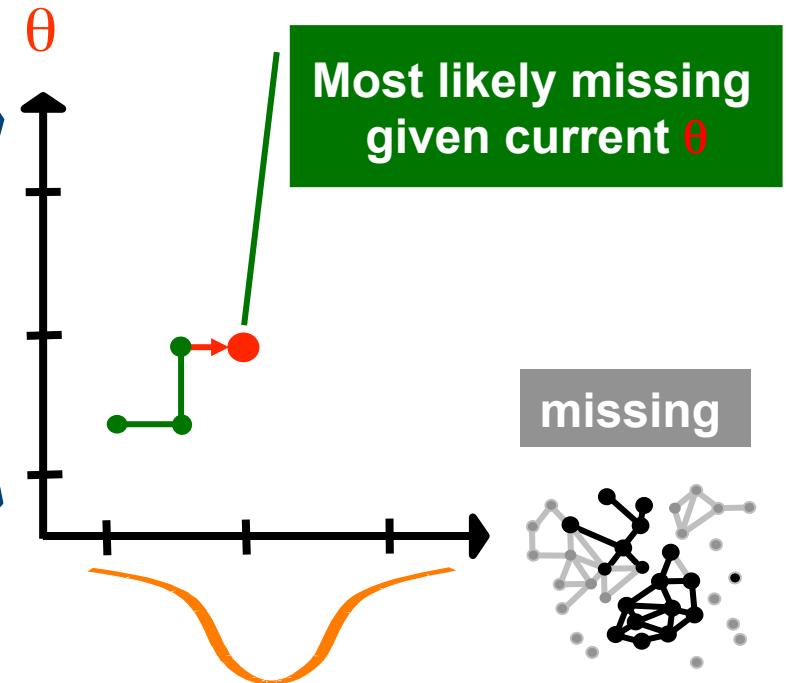
# Bayesian Data Augmentation

Simulate **parameters**



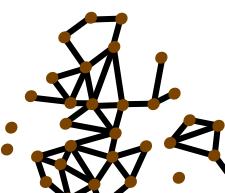
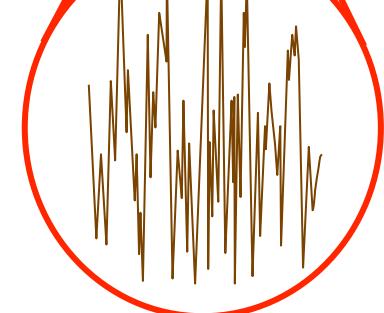
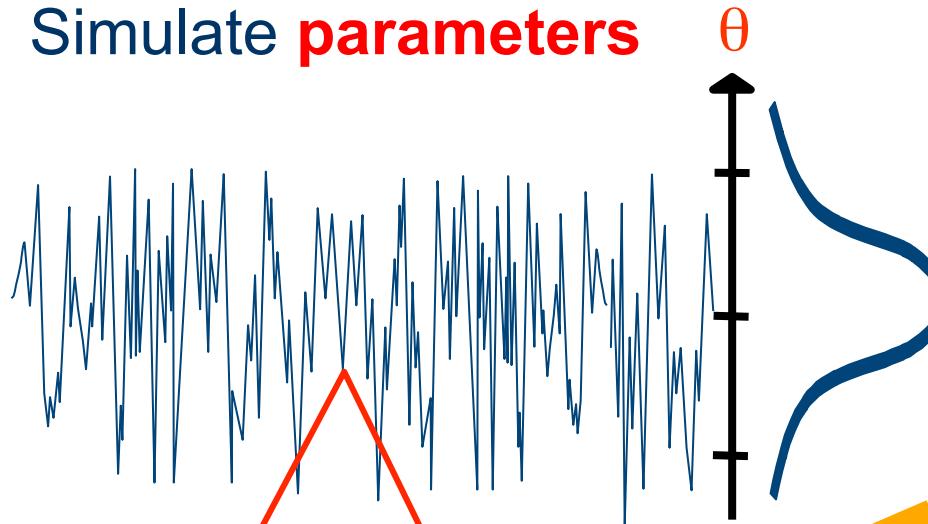
In each iteration  
simulate **graphs**

With missing data:



# Bayesian Data Augmentation

Simulate **parameters**



In each iteration  
simulate **graphs**

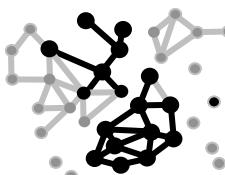
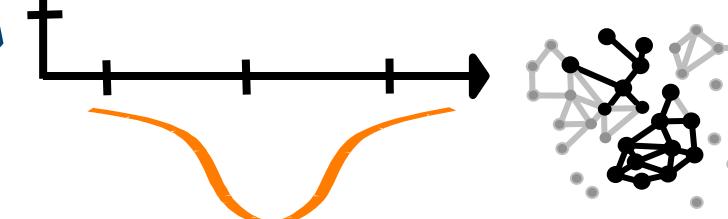
$\theta$

With missing data:

Most likely  $\theta$  given  
current missing

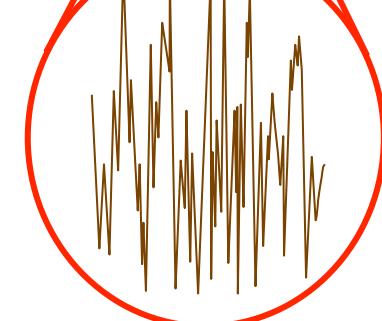
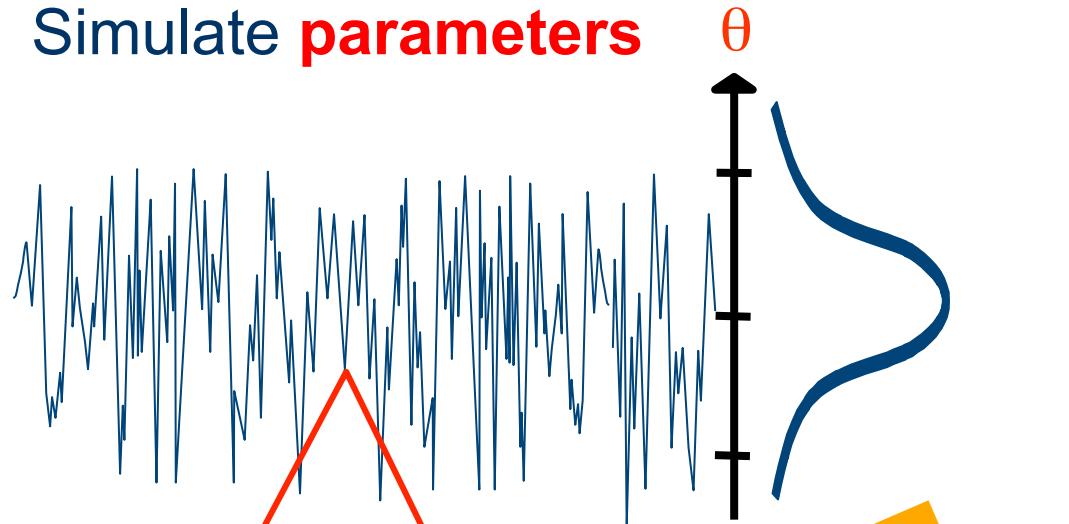
missing

$\theta$

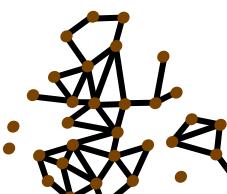


# Bayesian Data Augmentation

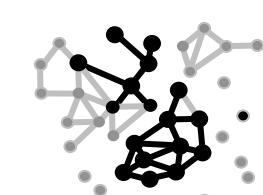
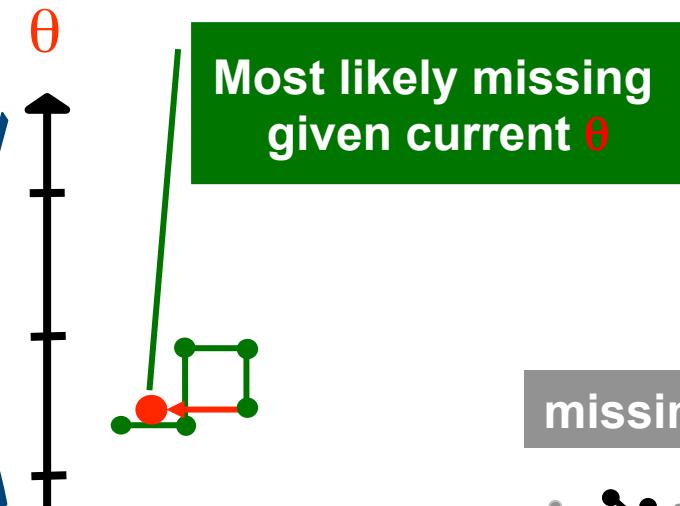
Simulate **parameters**



In each iteration  
simulate **graphs**

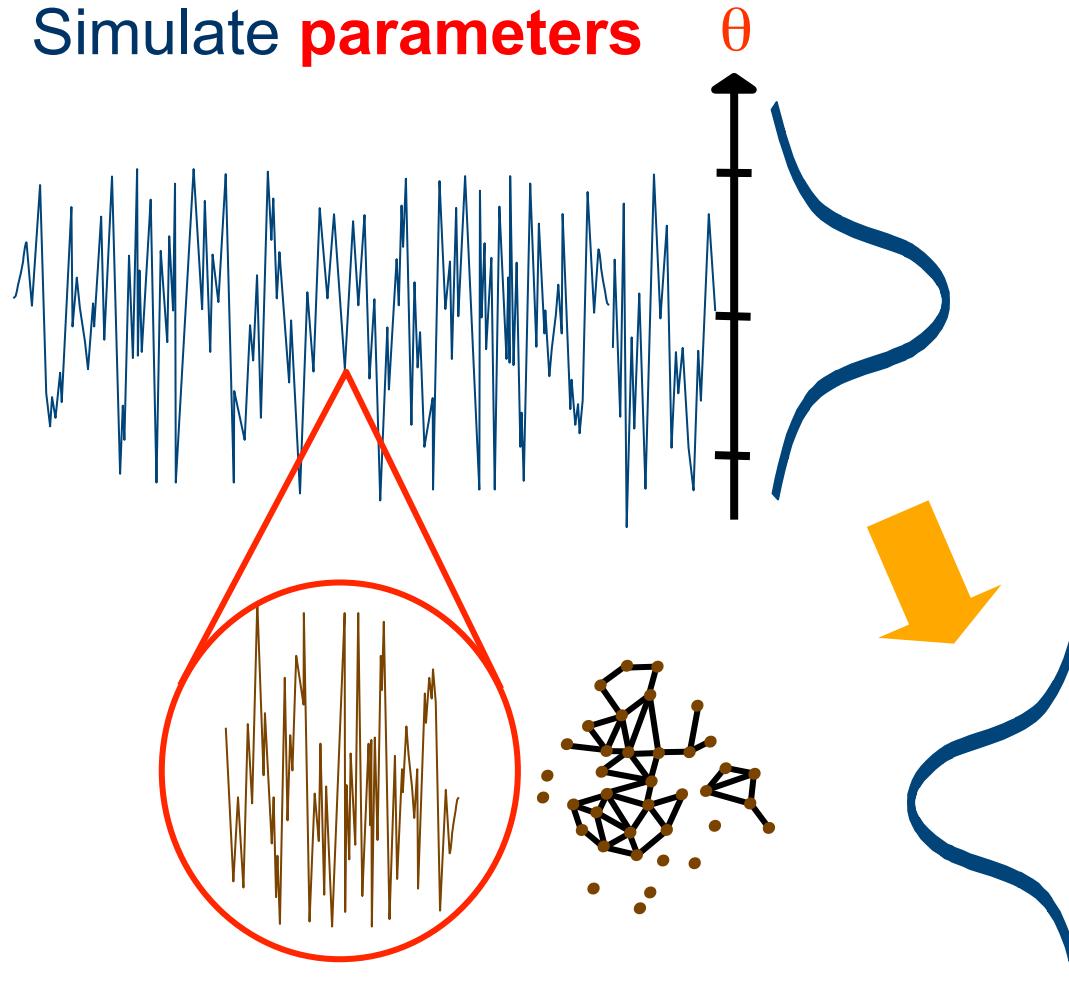


With missing data:

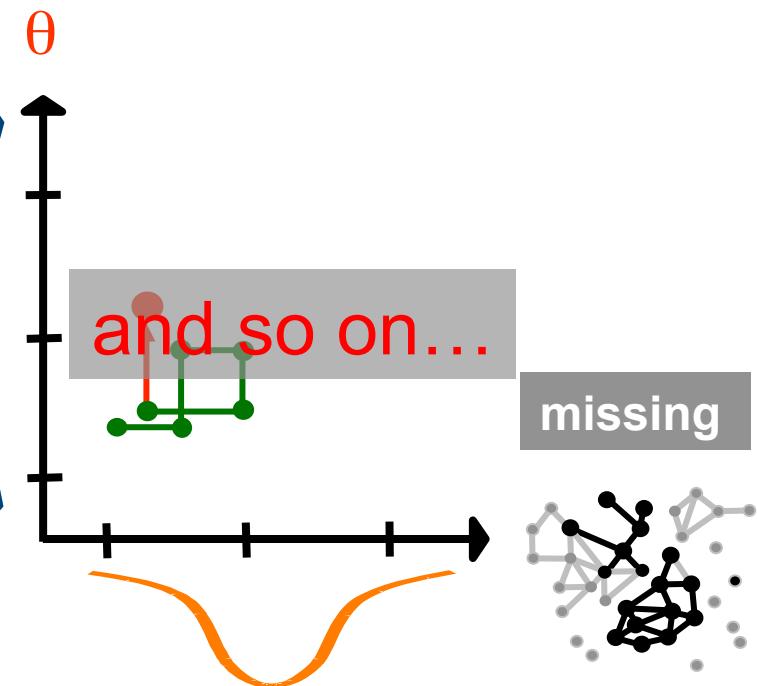


# Bayesian Data Augmentation

Simulate **parameters**



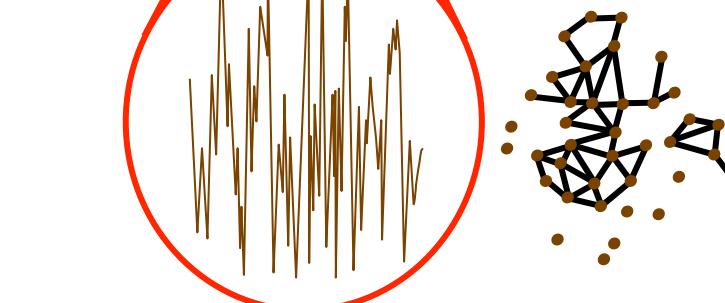
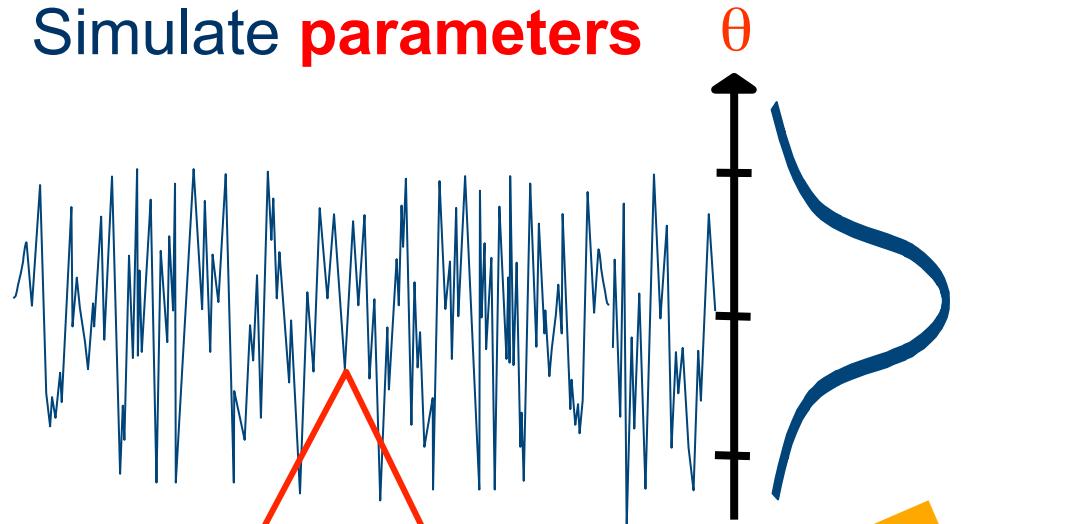
With missing data:



In each iteration  
simulate **graphs**

# Bayesian Data Augmentation

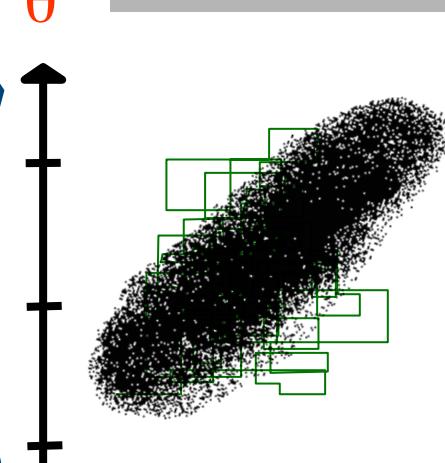
Simulate **parameters**



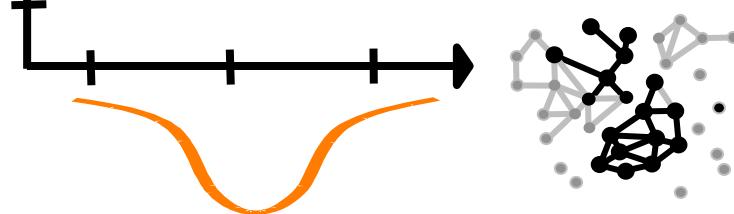
In each iteration  
simulate **graphs**

With missing data:

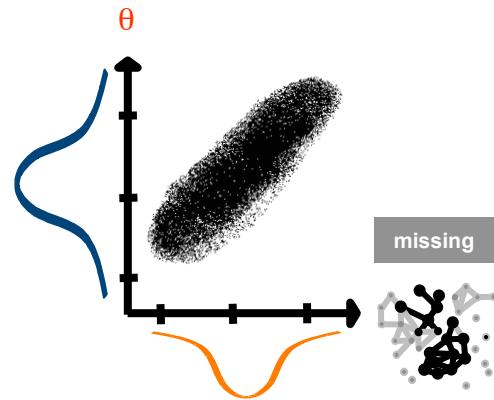
... until



missing



# Bayesian Data Augmentation



## What does it give us?

Distribution of parameters

Distribution of missing data

## Subtle point

Missing data does **not** depend on the parameters (we don't have to choose parameters to simulate missing)

# Lazega's (2001) Lawyers

Collaboration network among 36 lawyers in a  
New England law firm (Lazega, 2001)

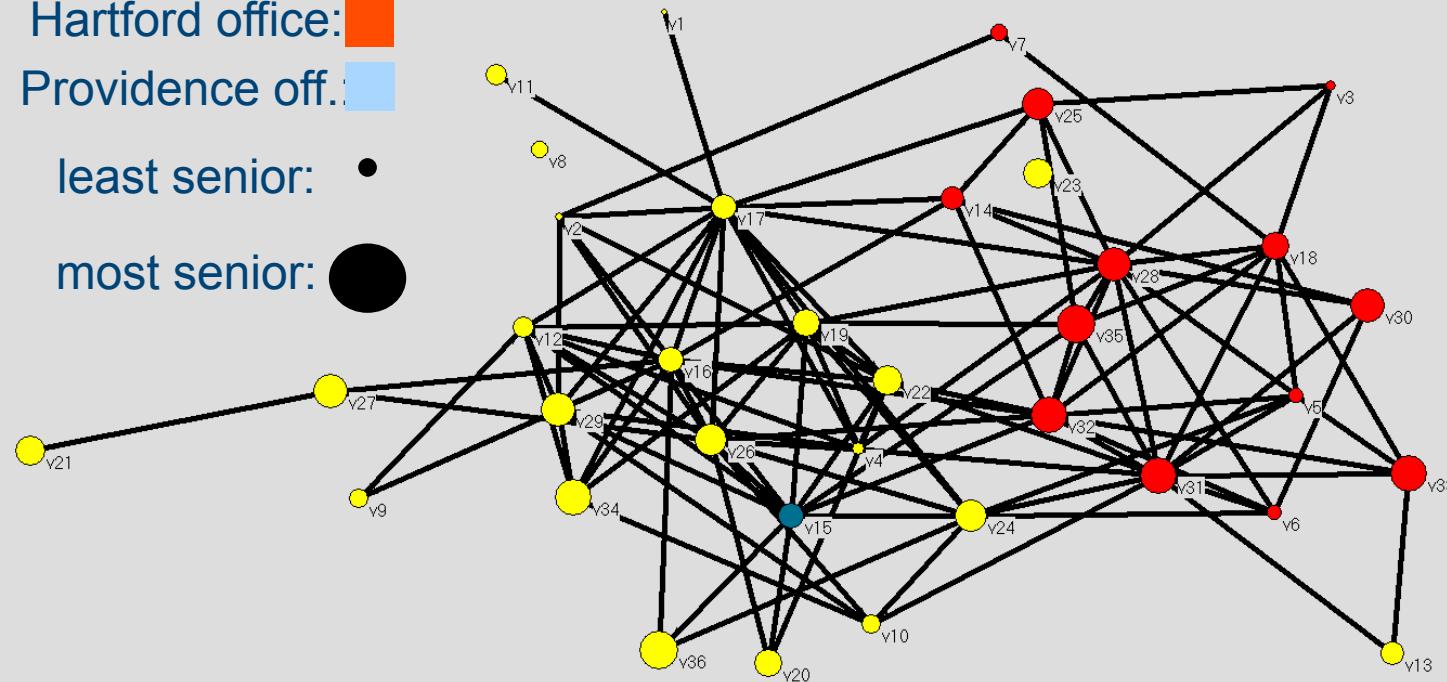
Boston office: ■

Hartford office: □

Providence off.: ▲

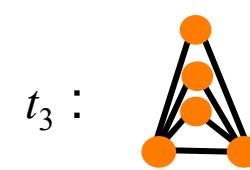
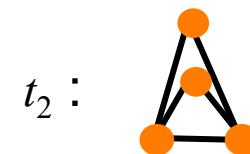
least senior: ●

most senior: ●



# Lazega's (2001) Lawyers

$(b_i = 1, \text{ if } i \text{ corporate},$   
 $0 \text{ litigation})$



etc.

**Main effect**

**Homophily**

**GWESP:**

**Edges:**

**Seniority:**

**Practice:**

**Practice:**

**Sex:**

**Office:**

$$\sum x_{ij}$$

$$\sum x_{ij}(a_i + a_j)$$

$$\sum x_{ij}(b_i + b_j)$$

$$\sum x_{ij} \mathbf{1}(b_i = b_j)$$

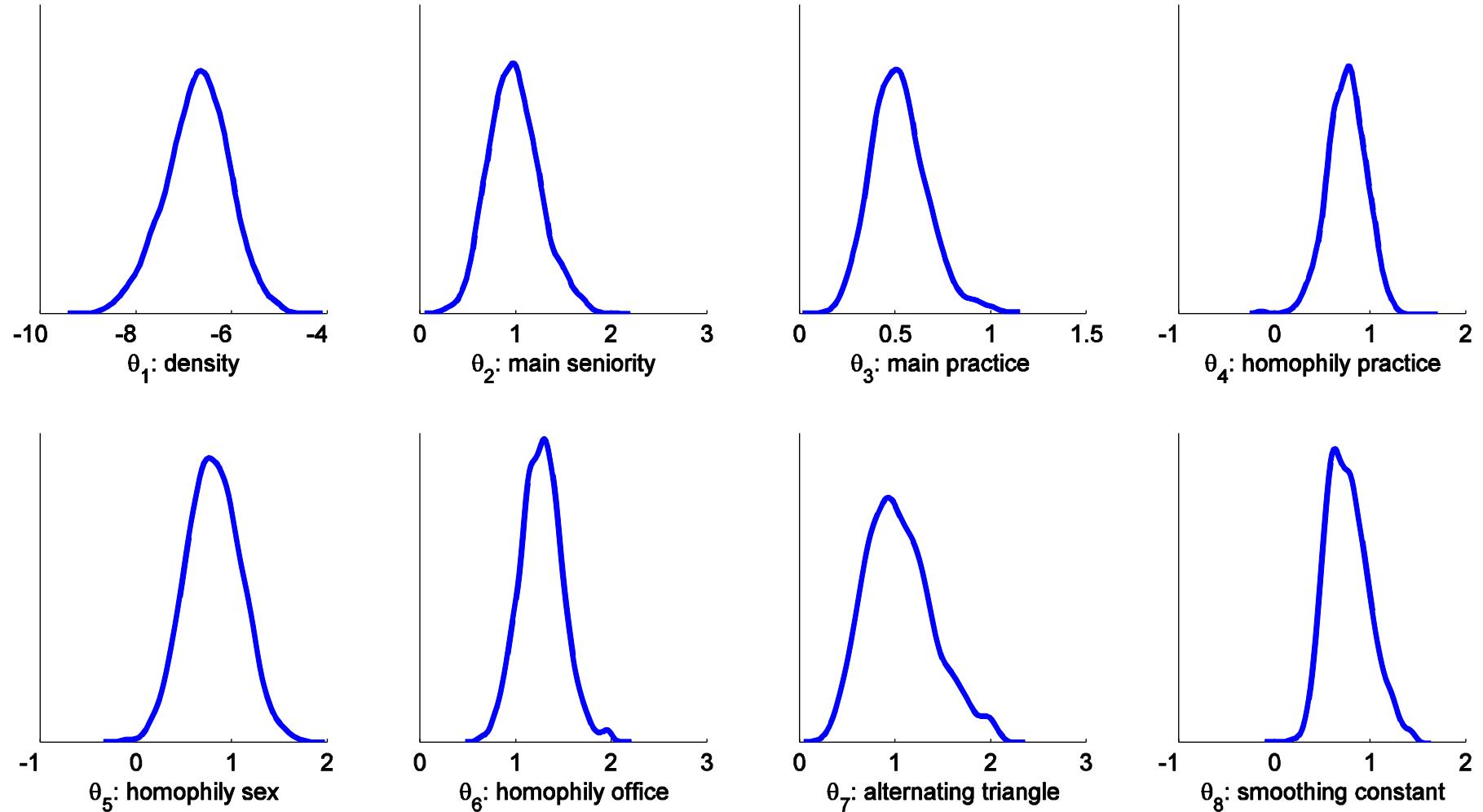
$$\sum x_{ij} \mathbf{1}(c_i = c_j)$$

$$\sum x_{ij} \mathbf{1}(d_i = d_j)$$

$$3t_1(x) - \frac{t_2(x)}{\lambda^1} + \dots + (-1)^{n-3} \frac{t_{n-2}(x)}{\lambda^{n-3}}$$

$$\text{with } \theta_8 = \log(\lambda)$$

# Lazega's (2001) Lawyers – ERGM posteriors (Koskinen, 2008)



Remove 200 of the 630 dyads at random

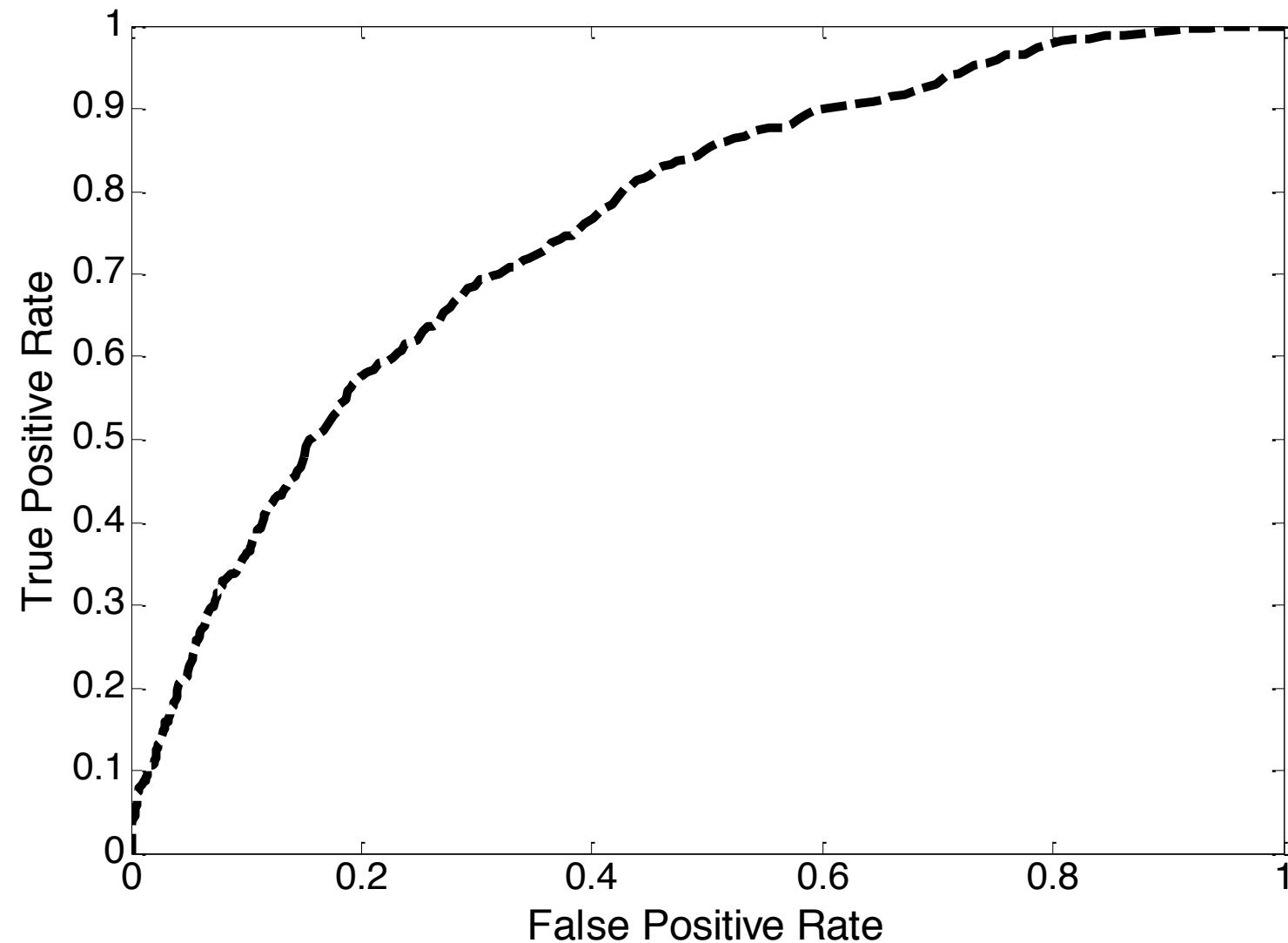
Fit inhomogeneous Bernoulli model obtain the posterior predictive tie-probabilities for the missing tie-variables

Fit ERGM and obtain the posterior predictive tie-probabilities for the missing tie-variables  
(Koskinen et al., in press)

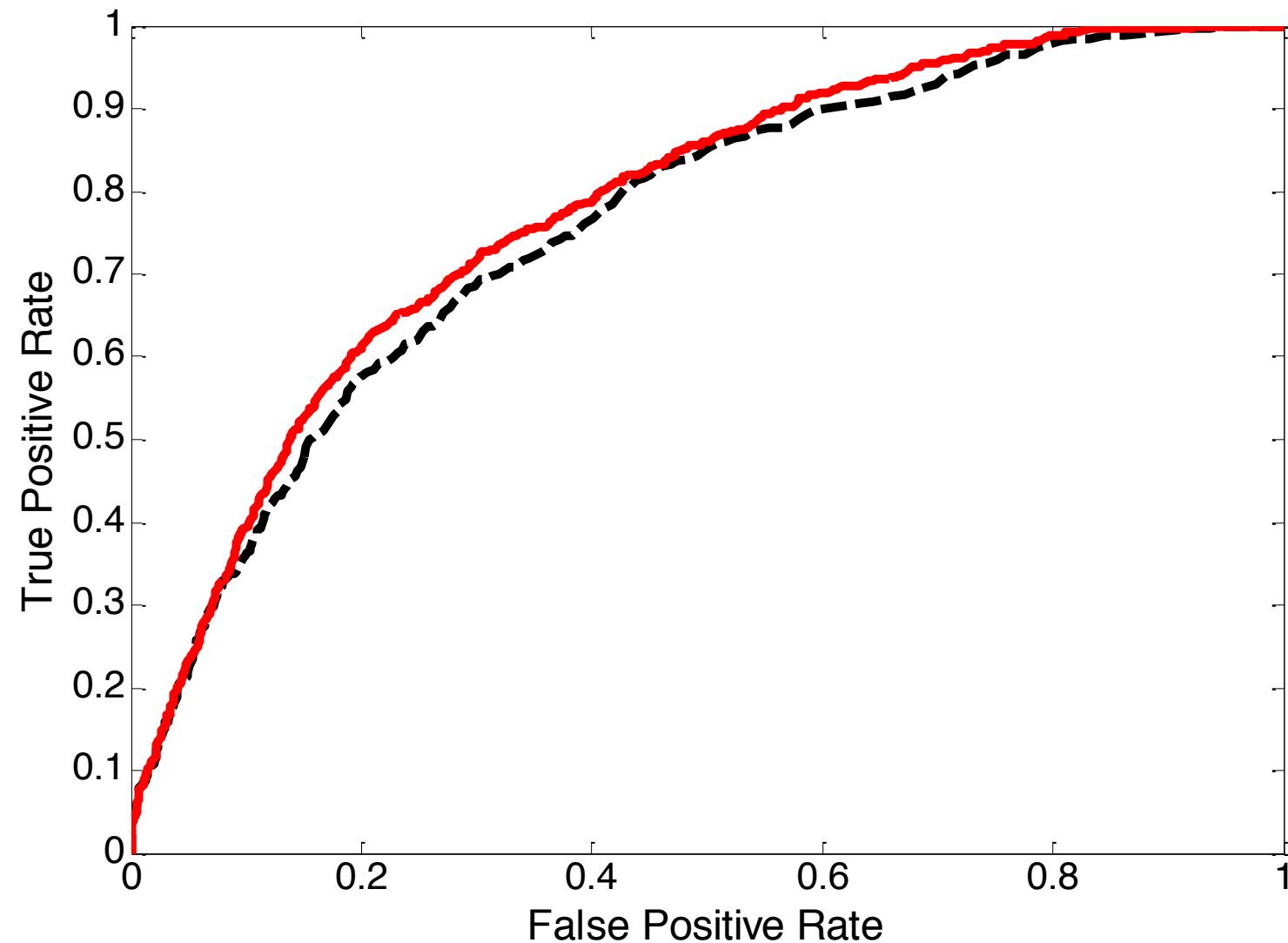
Fit Hoff's (2008) latent variable probit model with linear predictor  $\theta^T z(x_{ij}) + w_i \Lambda w_j^T$

Repeat many times

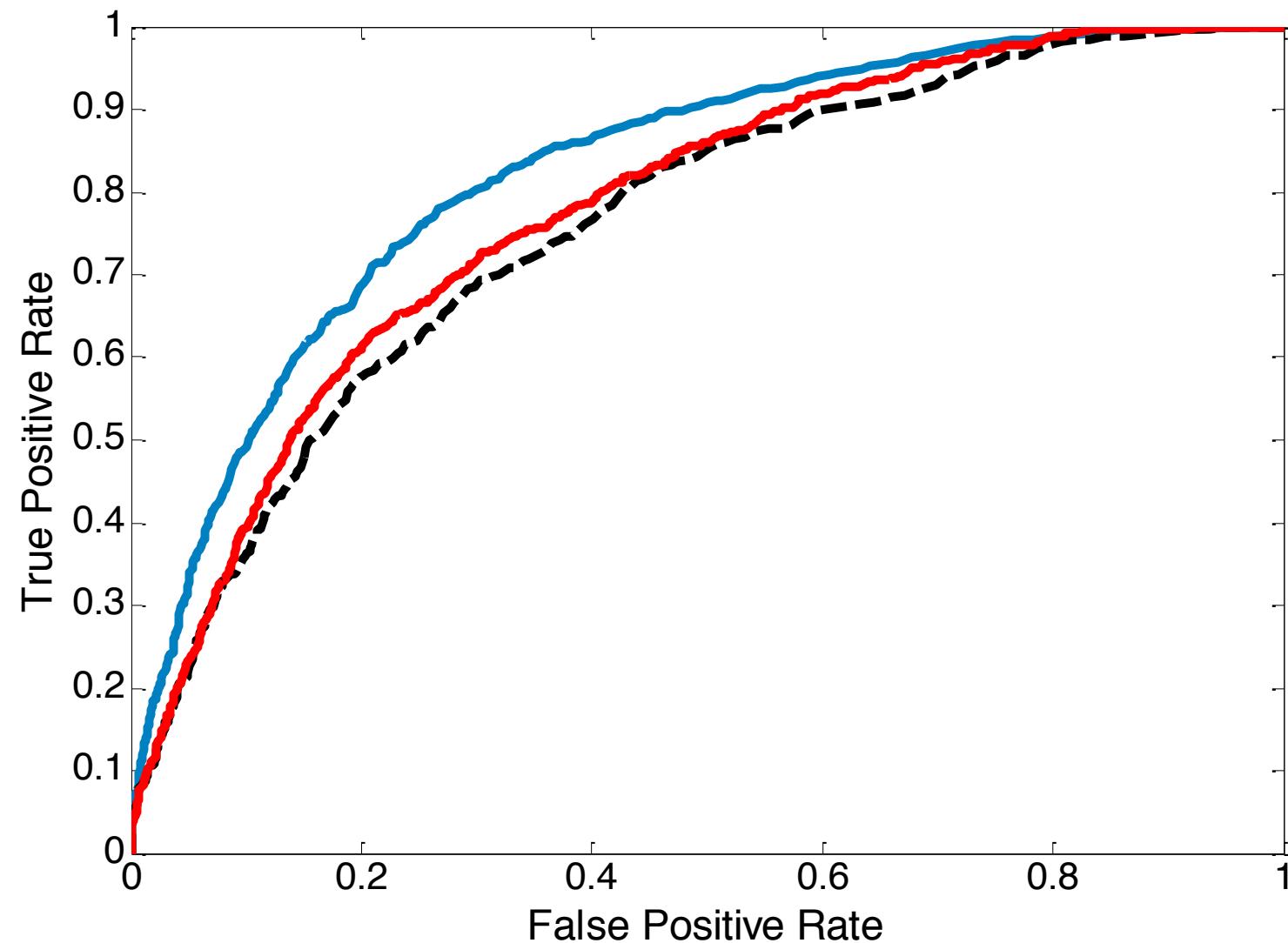
# ROC curve for predictive probabilities combined over 20 replications (Koskinen et al. 2010)



# ROC curve for predictive probabilities combined over 20 replications (Koskinen et al. 2010)



# ROC curve for predictive probabilities combined over 20 replications (Koskinen et al. 2010)



# Snowball sampling

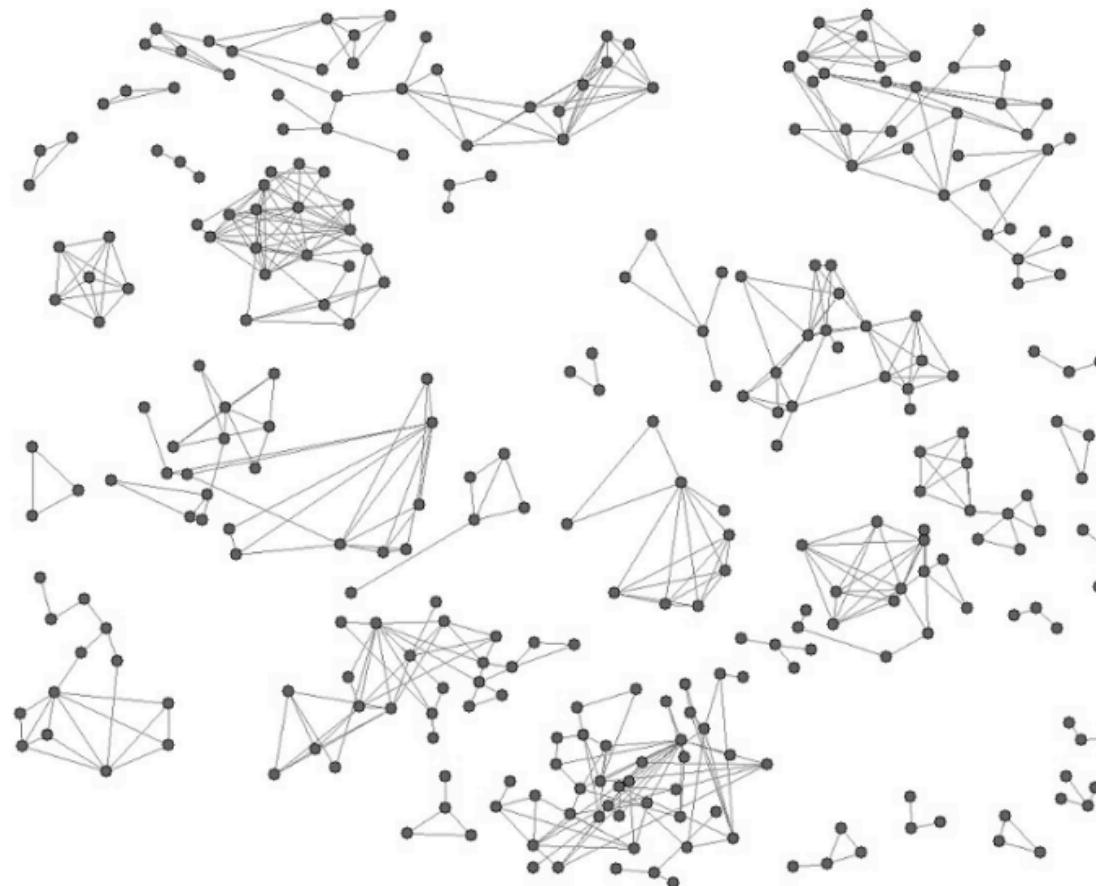
- Snowball sampling design ignorable for ERGM  
**(Thompson and Frank, 2000, Handcock & Gile 2010; Koskinen, Robins & Pattison, 2010)**
- ... **but** snowball sampling rarely used when population size is known...
- Using the Sageman (2004) “clandestine” network as test-bed for unknown  $N$

# Part 10

## Spatial embedding

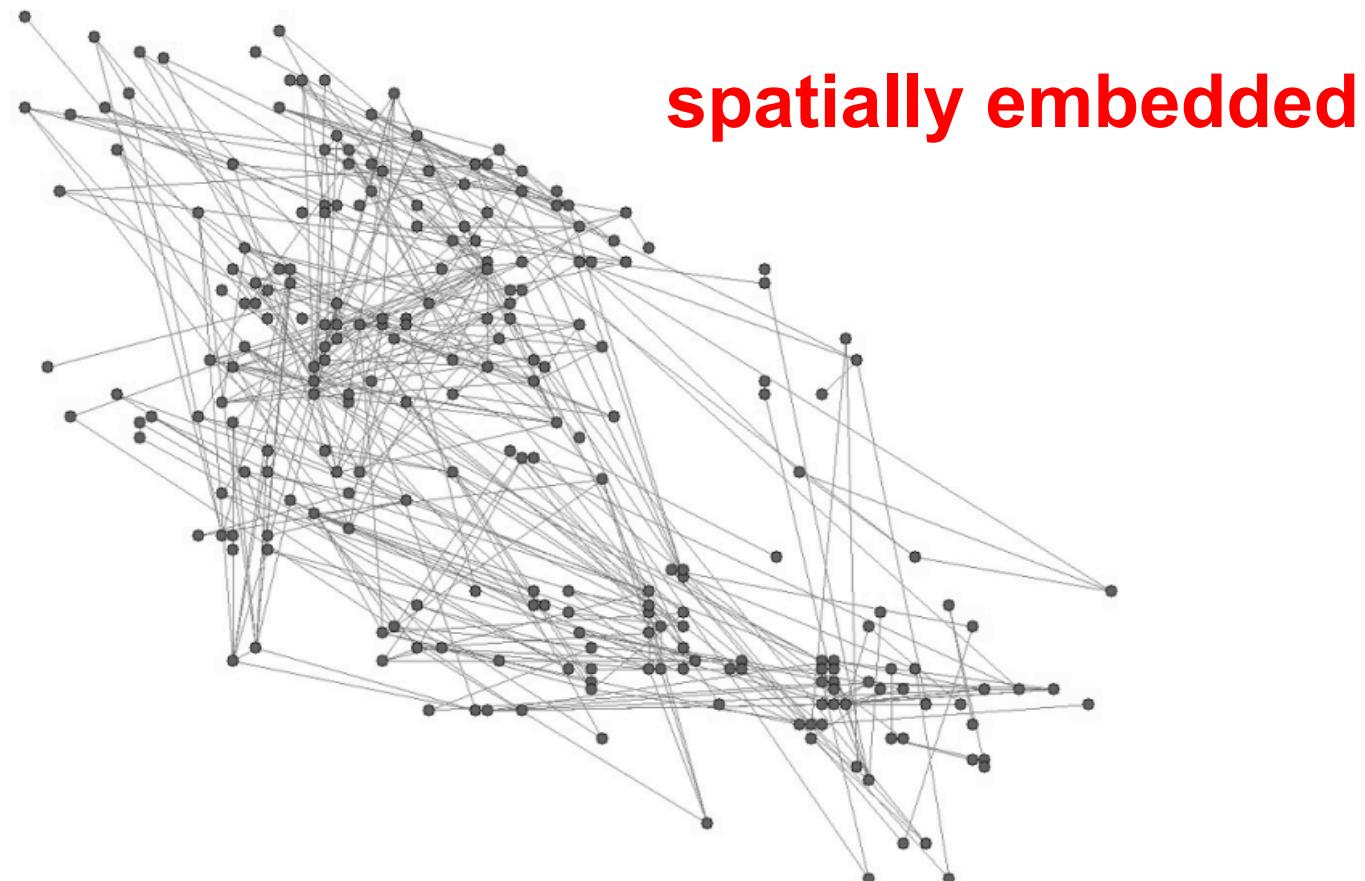
# Spatial embedding – Daraganova et al. 2011 SOCNET

306 actors in Victoria, Australia



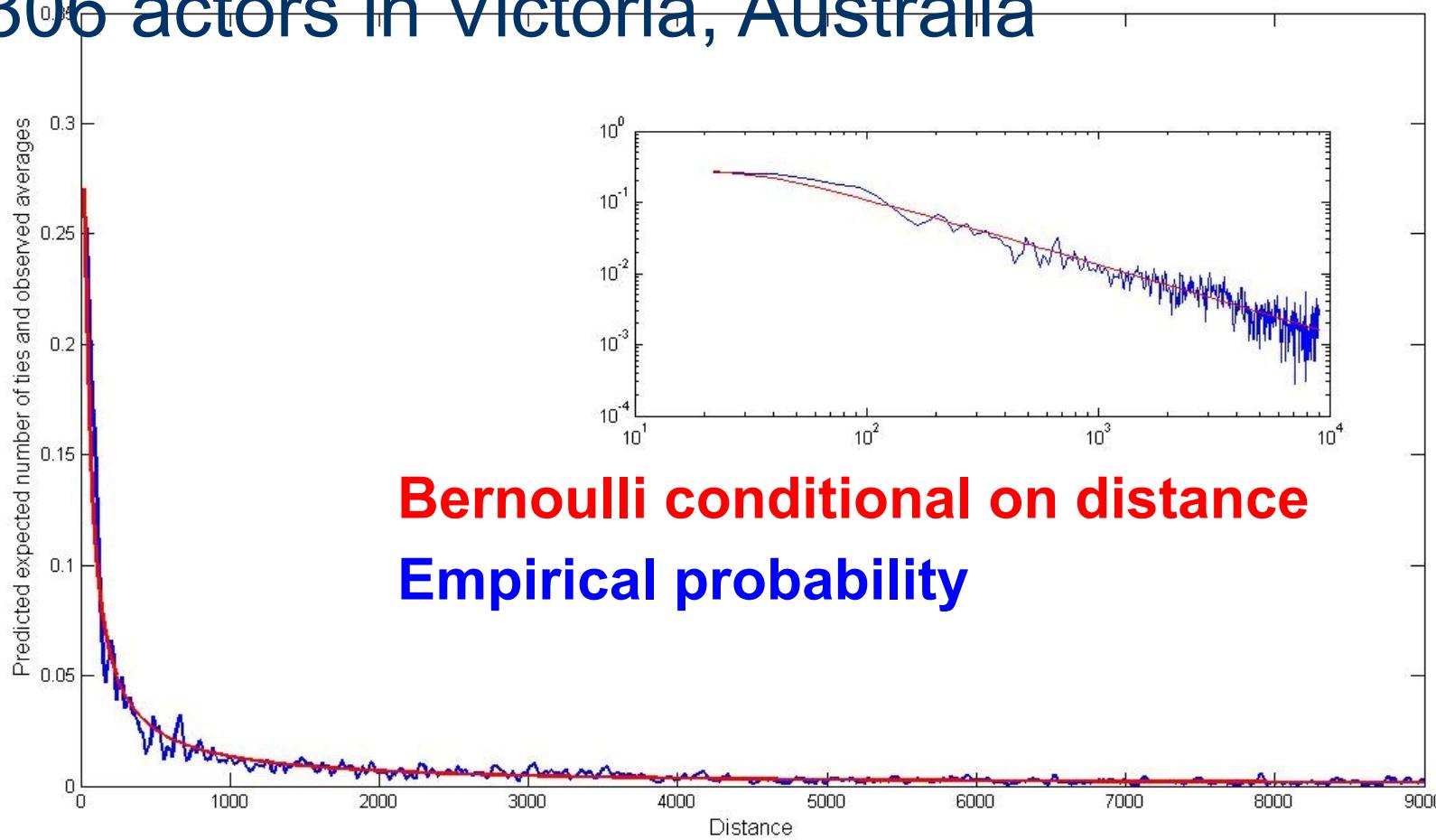
## Spatial embedding – Daraganova et al. 2011 SOCNET

306 actors in Victoria, Australia



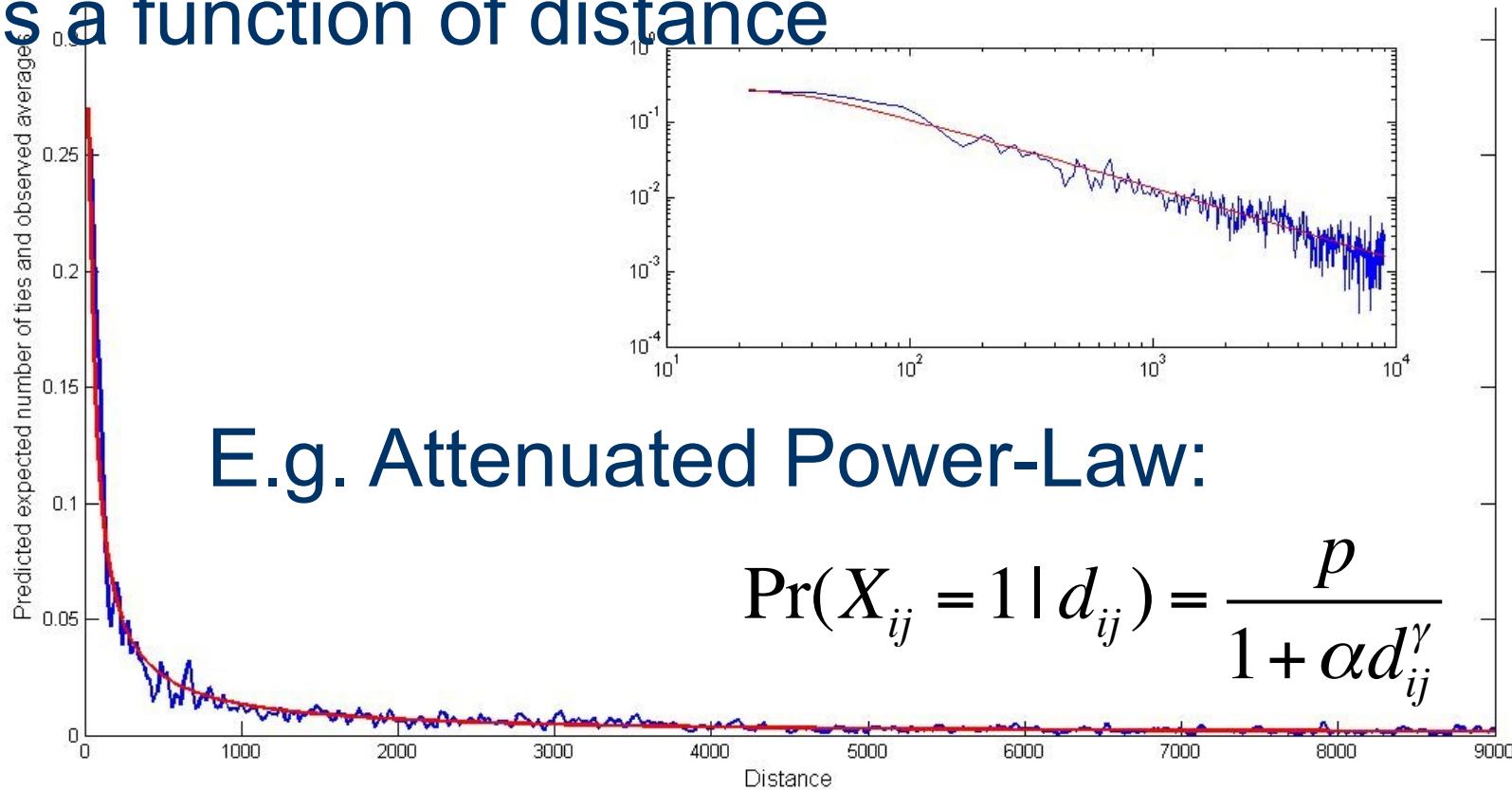
... all living within 14 kilometres of each other

306 actors in Victoria, Australia



... all living within 14 kilometres of each other

## Spatial interaction function: Tie probability as a function of distance



## Spatial interaction function: Tie probability as a function of distance

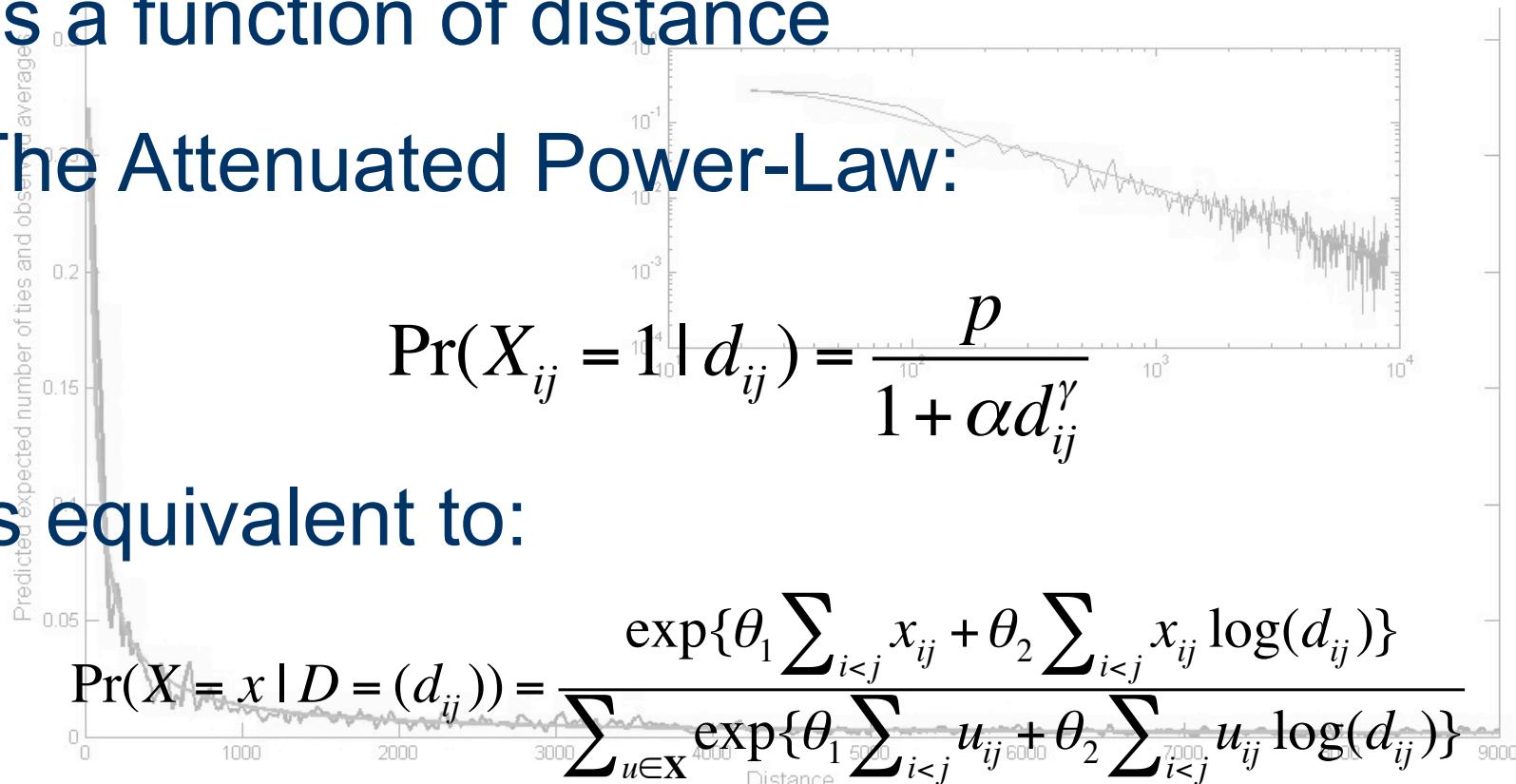
The Attenuated Power-Law:

$$\Pr(X_{ij} = 1 | d_{ij}) = \frac{p}{1 + \alpha d_{ij}^\gamma}$$

Is equivalent to:

$$\Pr(X = x | D = (d_{ij})) = \frac{\exp\{\theta_1 \sum_{i < j} x_{ij} + \theta_2 \sum_{i < j} x_{ij} \log(d_{ij})\}}{\sum_{u \in X} \exp\{\theta_1 \sum_{i < j} u_{ij} + \theta_2 \sum_{i < j} u_{ij} \log(d_{ij})\}}$$

with:  $p = 1$     $\alpha = e^{-\theta_1}$     $\gamma = -\theta_2$    AND:    $\log(d_{ij})$

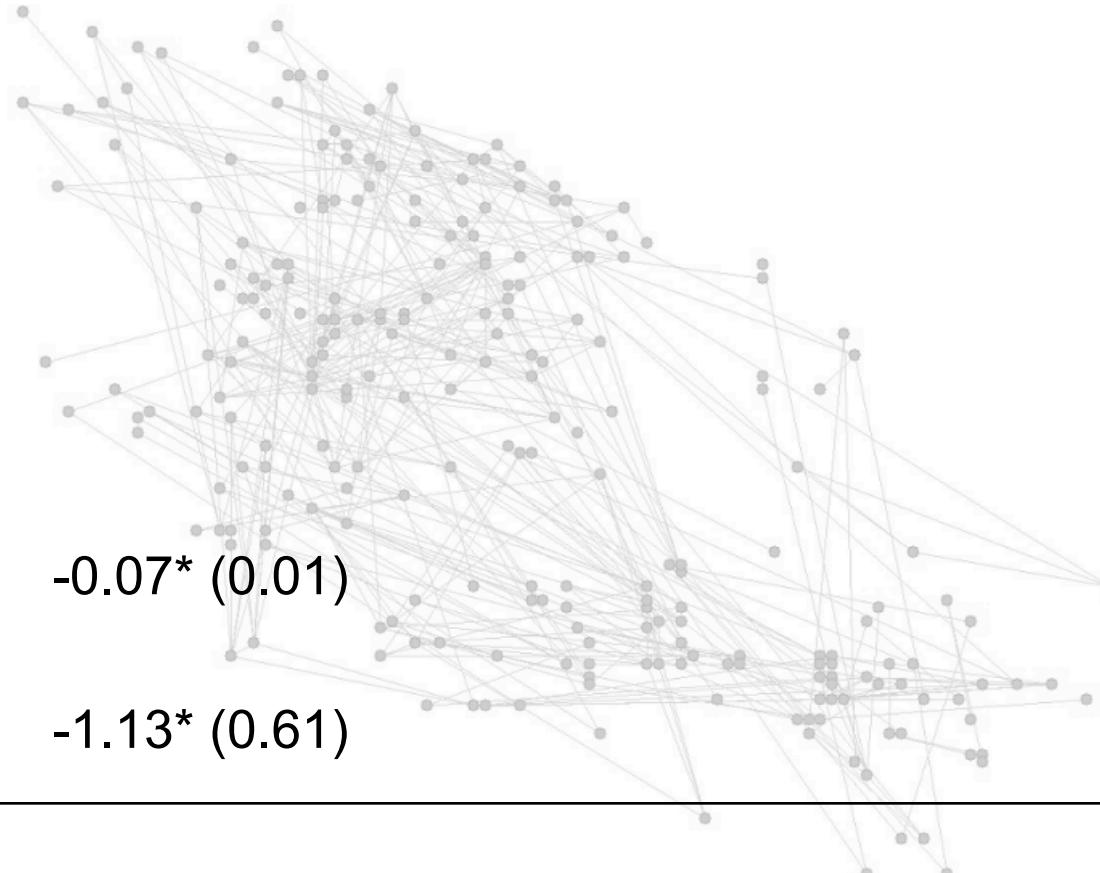


# Spatial embedding – Daraganova et al. 2011 SOCNET

Edges

-4.87\* (0.13)

Alt. star



Alt. triangel

Log distance

Age

heterophily

Gender

homophily

-0.07\* (0.01)

-1.13\* (0.61)

# Spatial embedding – Daraganova et al. 2011 SOCNET

Edges      -4.87\* (0.13)      1.56\* (0.65)

Alt. star

Alt. triangel

Log distance

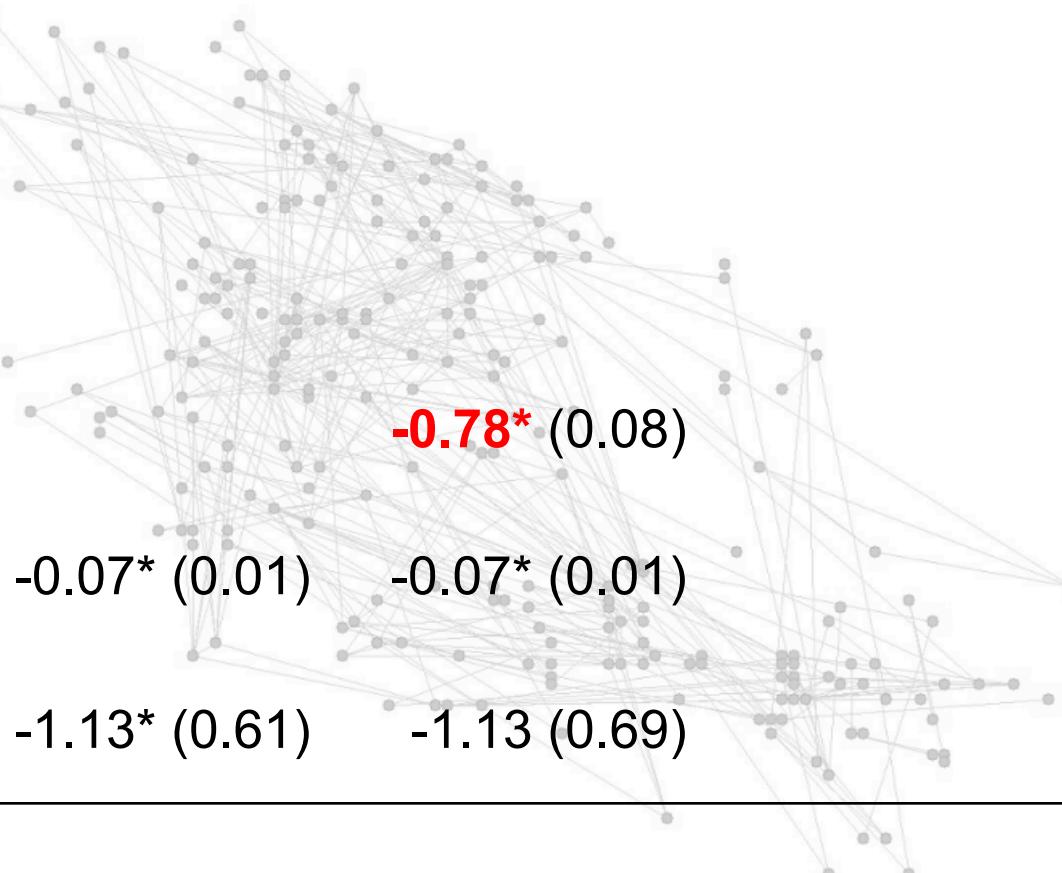
Age

heterophily

Gender

homophily

**-0.78\*** (0.08)



# Spatial embedding – Daraganova et al. 2011 SOCNET

Edges	-4.87* (0.13)	1.56* (0.65)	-4.79* (0.66)
-------	---------------	--------------	---------------

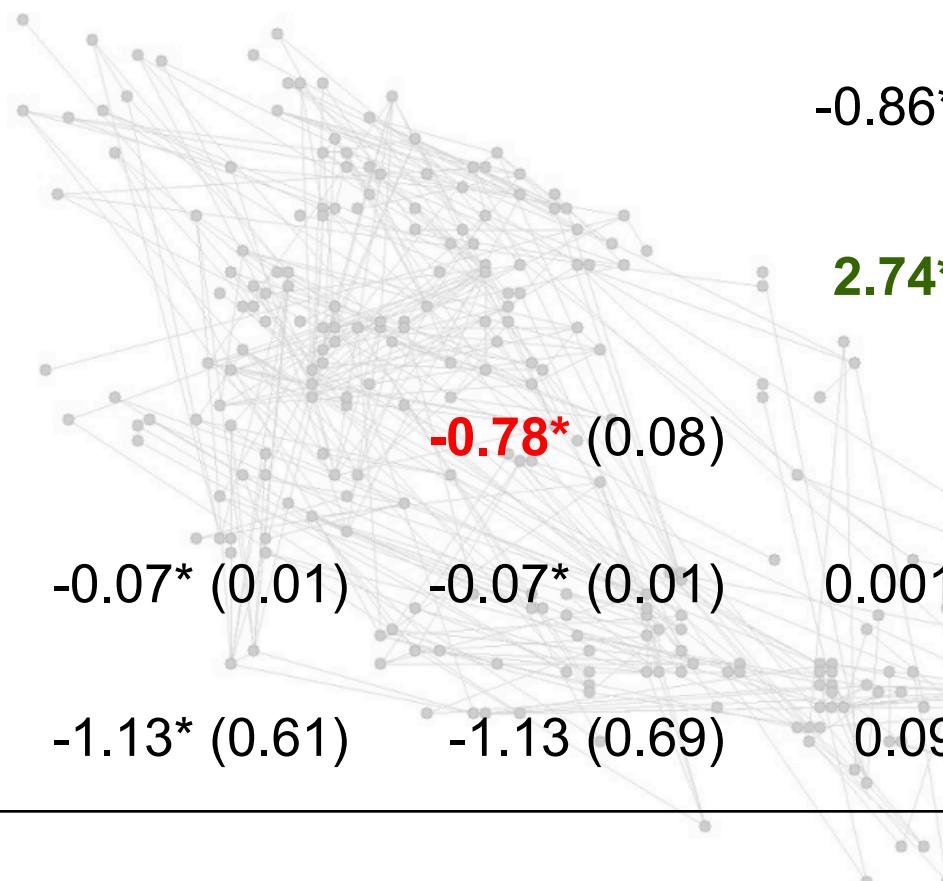
Alt. star		-0.86* (0.18)
-----------	--	---------------

Alt. triangel		<b>2.74*</b> (0.15)
---------------	--	---------------------

Log distance		<b>-0.78*</b> (0.08)
--------------	--	----------------------

Age heterophily	-0.07* (0.01)	-0.07* (0.01)	0.001 (0.07)
-----------------	---------------	---------------	--------------

Gender homophily	-1.13* (0.61)	-1.13 (0.69)	0.09 (0.83)
------------------	---------------	--------------	-------------



# Spatial embedding – Daraganova et al. 2011 SOCNET

Edges	-4.87* (0.13)	1.56* (0.65)	-4.79* (0.66)	-0.20 (0.87)
Alt. star			-0.86* (0.18)	-0.86* (0.2)
Alt. triangel			<b>2.74*</b> (0.15)	<b>2.69*</b> (0.14)
Log distance		<b>-0.78*</b> (0.08)		<b>-0.56*</b> (0.07)
Age heterophily	-0.07* (0.01)	-0.07* (0.01)	0.001 (0.07)	0.002 (0.06)
Gender homophily	-1.13* (0.61)	-1.13 (0.69)	0.09 (0.83)	0.07 (0.47)

**ERGM: *distance* and *endogenous* dependence explain different things**

# Part 8

## Further issues

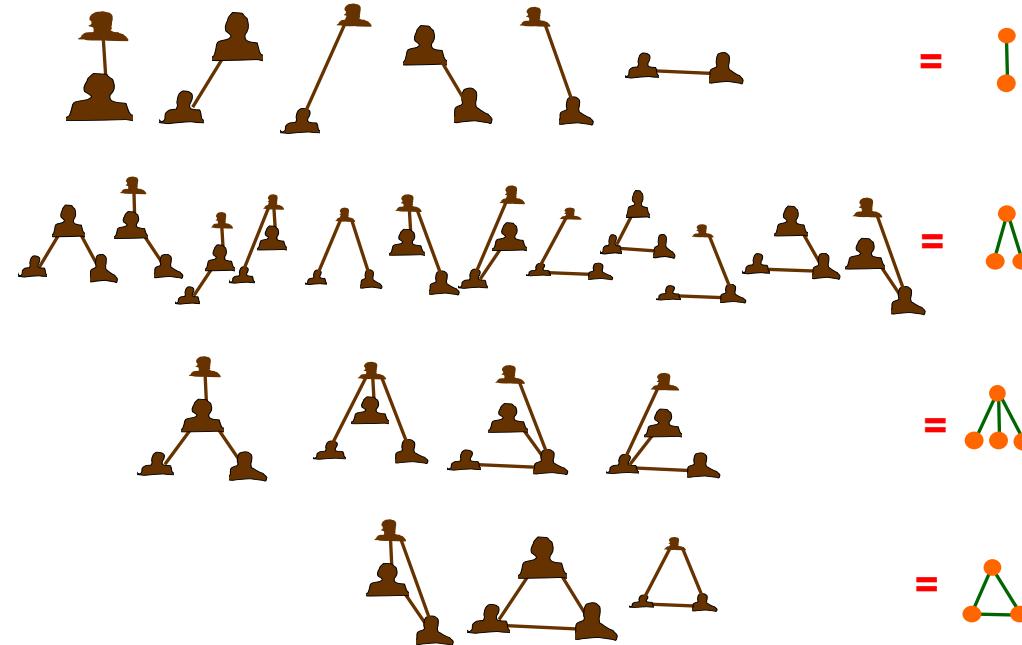
## Other extensions

There are **ERGMs** (ERGM-like models) for

- directed data
- valued data
- bipartite data
- multiplex data
- longitudinal data
- modelling actor autoregressive attributes

## Further issues – relaxing homogeneity assumption

ERGMs typically assume homogeneity



(A) Block modelling and ERGM (Koskinen, 2009)

(B) Latent class ERGM (Schweingberger & Handcock)

### Assessing Goodness of Fit:

- Posterior predictive distributions (Koskinen, 2008; Koskinen, Robins & Pattison, 2010; Caimo & Friel, 2011)
- Non-Bayesian heuristic GOF (Robins et al., 2007; Hunter et al., 2008; Robins et al., 2009; Wang et al., 2009)

### Model selection

- Path sampling for AIC (Hunter & Handcock, 2006); Conceptual **caveat**: model complexity when variables dependent?
- Bayes factors (Wang & Handcock...?)

## ERGMs

- Increasingly being used
- Increasingly being understood
- Increasingly being able to handle imperfect data (also missing link prediction)

## Methods

- Plenty of open issues
- Bayes is the way of the future

## Legitimacy and dissemination

- e.g. Lusher, Koskinen, Robins ERGMs for SN, CUP, 2011

# Remaining question: used to be p\*...

... why ERGM?

