Chaos, Complexity, and Inference (36-462) Lecture 21

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Models of Networks, with Origin Myths

Erdős-Rényi Encore

Erdős-Rényi with Node Types

Watts-Strogatz "Small World" Graphs

Exponential-Family Random Graphs

Preferential Attachment

Erdős-Rényi Again

n nodes, edges are IID binary variables with probability p Degree of node $i = K_i$

$$K_i \sim \text{Binom}(n-1,p) \rightsquigarrow \text{Pois}(np)$$

Problems

Degree distribution Not Poisson

Reciprocity
$$Pr(A_{jj} = 1 | A_{ij} = 1) > p$$

Transitivity
$$Pr(A_{ik} = 1 | A_{ij} = A_{jk} = 1) > p$$

Homophily/Assortativeness
$$Pr(A_{ij} = 1 | type_i = type_j) > p$$



Inhomogeneous E-R Models

Give each node a type, $1, \ldots k, T_i$

mixing matrix P_{ab} = probability of link from type a to type b

Edges are still independent given type

Edges are *not* independent ignoring type

Example: k = 2, types uniform and independent

$$P = \left[\begin{array}{cc} 0.9 & 0.1 \\ 0.1 & 0.9 \end{array} \right]$$

Obviously gives homophily

$$p = \Pr(A_{ij} = 1)$$

$$= 0.9 \Pr(T_i = T_j = 1) + 0.1 \Pr(T_i = 1, T_j = 2)$$

$$+0.1 \Pr(T_i = 2, T_j = 1) + 0.9 \Pr(T_i = T_j = 2)$$

$$= 0.9 \times 0.25 + 0.1 \times 0.25 + 0.1 \times 0.25 + 0.9 \times 0.25 = 0.5$$

Also gives reciprocity:

$$\begin{aligned} &\Pr\left(A_{ji=1}=1,A_{ij}=1\right)\\ &= &0.81\Pr\left(T_{i}=T_{j}=1\right)+0.01\Pr\left(T_{i}=1,T_{j}=2\right)\\ &+0.01\Pr\left(T_{i}=2,T_{j}=1\right)+0.81\Pr\left(T_{i}=T_{j}=2\right)\\ &= &0.41\\ &\Pr\left(A_{ji=1}=1|A_{ij}=1\right)\\ &= &\frac{\Pr\left(A_{ji}=1,A_{ij}=1\right)}{\Pr\left(A_{ij}=1\right)}\\ &= &0.82>0.5 \end{aligned}$$

EXERCISE: Show that this model has transitivity of edges as well

One direction for extending this: **block models** ("block" = type), indicating "type A gets links from type B, gives links to type C, never gets links from D or E..."

Community structure or **modularity** is a limiting case of this, where mixing matrix has big diagonal entries, small off-diagonal ones

References: Reichardt and White (2007) for discovering block models; Clauset *et al.* (2007) for discovering hierarchies of modules; http://bactra.org/notebooks/community-discovery.html for references on community structure and community discovery

Watts-Strogatz "Small World" Graphs

Watts and Strogatz (1998)

Regular lattices have a lot of reciprocity and transitivity/clustering

but are "large worlds", in d dimensions diameter

$$= O(n^{1/d}) \gg O(\log n)$$

Somehow interpolate between lattices and E-R graphs to get all three properties

but work with undirected graphs for simplicity

Solution: start with regular lattice, add "long-range shortcuts" at random

First approach: For each edge, with probability ρ , re-wire one edge to a uniformly random new node (avoiding self-loops)

As $\rho \rightarrow$ 0, go to regular lattice

As $\rho \to 1$, go to E-R graph with same density as lattice can create disconnected graphs

Second approach: add random edges *without* removing old ones

easier to manipulate, doesn't quite go to E-R as $\rho \to 1$ Will do more with this in the EXERCISES



Exponential Family Random Graphs

Measure graph properties like density, reciprocity, transitivity; specify graph probabilities in terms of them **Exponential families** are the easiest way to do this

$$\Pr(X = x) = \frac{h(x) \exp\left\{\sum_{i=1}^{d} \theta_i T_i(x)\right\}}{\int dx \ h(x) \exp\left\{\sum_{i=1}^{d} \theta_i T_i(x)\right\}}$$
$$= \frac{h(x) \exp\left\{\sum_{i=1}^{d} \theta_i T_i(x)\right\}}{Z(\theta)}$$

 T_i are **sufficient statistics**, θ_i are **natural parameters** Acronym: ERGM, Exponential family Random Graph Model ("err-gim" or

"err-gum")



E-R model is an exponential family:

$$\Pr(A = a) = \prod_{i=1}^{n} \prod_{j \neq i} p^{a_{ij}} (1 - p)^{(1 - a_{ij})}$$

$$= p^{\sum_{ij} a_{ij}} (1 - p)^{n(n-1) - \sum_{ij} a_{ij}}$$

$$= (1 - p)^{n(n-1)} \left(\frac{p}{1 - p}\right)^{\sum_{ij} a_{ij}}$$

$$= (1 - p)^{n(n-1)} \exp\left\{ (\log p/(1 - p)) \sum_{ij} a_{ij} \right\}$$

so
$$T = \sum_{ij} a_{ij}$$
, $\theta = \log p/(1-p)$, $Z(\theta) = (1-p)^{-n(n-1)}$



Exponential family models are easy to fit by maximum likelihood, if you can find $Z(\theta)$ or $\mathbf{E}_{\theta}[T_i(x)]$

$$\frac{\partial \log \Pr(X = x)}{\partial \theta_{i}}$$

$$= \frac{\partial}{\partial \theta_{i}} \log h(x) + \frac{\partial}{\partial \theta_{i}} \sum_{j=1}^{d} \theta_{j} T_{j}(x) - \frac{\partial}{\partial \theta_{i}} \log Z(\theta)$$

$$= 0 + T_{i}(x) - \frac{1}{Z(\theta)} \frac{\partial Z(\theta)}{\partial \theta_{i}}$$

$$= T_{i}(x) - \mathbf{E}_{\theta} [T_{i}(X)]$$

For E-R model,
$$\mathbf{E}_{\theta}\left[\sum_{ij}A_{ij}\right]=n(n-1)p$$
 so
$$\widehat{p}_{MLE}=\frac{\sum_{ij}a_{ij}}{n(n-1)}$$

What about more complicated ERGMs?

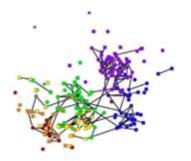
"p₁ model": sufficient statistics are total number of edges, and total number of *reciprocal* edges

Not so easy to solve but can be done (Wasserman and Faust, 1994; Hunter *et al.*, 2008)

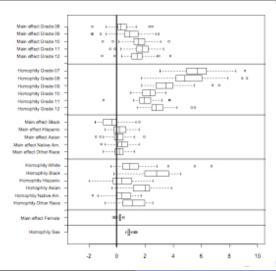
*p**: general ERGM, can add more features, homophily as such vs. reciprocity or transitivity as such...

Example of ERGMs Working

High school friendship network (Goodreau et al., 2005)



Fit model including homophily by sex, grade, race; also different over all probability of forming edges ("main effect")



Best R package: statnet (on CRAN)

Generally *not* possible to solve

Use simulation to approximate $Z(\theta)$ and/or $\mathbf{E}_{\theta}[T(X)]$ (Hunter and Handcock, 2006)

even then there can be pathologies from bad choice of model (e.g. model say probability of these network statistics is 10^{-50})

Some Important Weaknesses of ERGMs

- Possible pathologies in fitting
- ② "Statistics convenient for us to measure" ≠ "important causal variables"
- Matching some statistics doesn't mean matching others (Hunter et al., 2008)
- No origin myth/generative model (typically)

Some Generative Models

- E-R model edges appear and disappear independently *over time* (works whether or not homogeneous)
 - p₁ model Markov chain, edge in one direction makes adding edge more likely, losing one edge makes other tend to go away
- Watts-Strogatz Models See Clauset and Moore (2003) for a semi-plausible story about adaptive re-wiring
 - E-R again Add nodes one by one, each node adds links to existing nodes independently with probability *p*
- Preferential attachment Graphical version of Yule-Simon process



Preferential Attachment

Made famous by Barabási and Albert (1999); Albert and Barabási (2002)

At each time-step a new node arrives

With probability ρ , new node i makes edge to old node j,

picking $j \propto k_i$, degree of j

With probability $1 - \rho$, *i* links to a completely random node

This is *exactly* the Yule-Simon process that produces power law tails (Bornholdt and Ebel, 2001)

Apparently first applied to networks by Price (1965)

Will see more in the EXERCISES

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