# Algebraic Statistics of $p_{1}$ Network Models: Markov Bases and Their Uses 

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- $p_{1}$ model, log-linear model, and algebraic statistics:

Using algebra-geometry structure to "understand" a family of network models and frame a related set of statistical estimation and inference issues.

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- Can have multiple relationships; attributes for nodes and/or edges
- Edges can be directed or undirected
- Data available to be observed at time $t=t_{0}$
- Underlying stochastic process that describes the network structure and evolution.
- Come to my talk on dynamic network modeling this afternoon in Grpahical models Workshop!


## Example: The Framingham "Obesity" Study

- Original Framingham "sample" cohort begun in 1947.
- Offspring cohort of $N_{0}=5,124$ individuals measured beginning in 1971 for $T=7$ epochs centered at 1971, 1981, 1985, 1989, 1992, 1997, 1999.
- Link information on family members and one close friend.
- Total number of individuals on whom we have obesity measures is $\mathrm{N}=12,067$.
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Figure 1. Largest Connected Subcomponent of the Social Network in the Framingham Heart Study in the Year 2000.
Each circle (node) represents one person in the data set. There are 2200 persons in this subcomponent of the social network. Circles with red borders denote women, and circles with blue borders denote men. The size of each circle is proportional to the person's body-mass index. The interior color of the circles indicates the person's obesity status: yellow denotes an obese person (body-mass index,$\geq 30$ ) and green denotes a nonobese person. The colors of the ties between the nodes indicate the relationship between them: purple denotes a friendship or marital tie and orange denotes a familial tie.

## Example -The Collective Dynamics of Smoking in a Large Social Network (J.Fowler)

## Example: Monks in a Monastery

- 18 novices observed over two years.
- Network data gather at 4 time points; and on multiple relationships..
- See analyses in Airoldi, et al., (2008)


Negative praise


## Holland and Leinhardt's $p_{1}$ model

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\log \operatorname{Prob}(\text { no edge })=\log P_{i j}(0,0)=\lambda_{i j}
$$

$$
\log \operatorname{Prob}(\text { from } i \text { to } j)=\log P_{i j}(1,0)=\lambda_{i j}+\alpha_{i}+\beta_{j}+\theta
$$

$$
\log \operatorname{Prob}(\text { from } j \text { to } i)=\log P_{i j}(0,1)=\lambda_{i j}+\alpha_{j}+\beta_{i}+\theta
$$

$$
\log \operatorname{Prob}(\text { bi-directed edge })=\log P_{i j}(1,1)=\lambda_{i j}+\alpha_{i}+\beta_{j}+\alpha_{j}+\beta_{i}+2 \theta+\rho_{i j}
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\end{aligned}
$$

- 3 common forms:

$$
\begin{aligned}
& \rho_{i j}=0 \text { (no reciprocal effect) } \\
& \rho_{i j}=\rho \text { (constant reciprocation factor) } \\
& \rho_{i j}=\rho+\rho_{i}+\rho_{j} \text { (edge-dependent reciprocation) }
\end{aligned}
$$

## Estimation for $p_{1}$

- The likelihood function for the $p_{1}$ model is clearly of exponential family form.
- For the constant reciprocation version, we have

$$
\begin{equation*}
\log p_{1}(x) \propto x_{++} \theta+\sum_{i} x_{i+} \alpha_{i}+\sum_{j} x_{+} j \beta_{j}+\sum_{i j} x_{i j} x_{j i} \rho \tag{1}
\end{equation*}
$$

- Holland-Leinhardt explored goodness of fit of model empirically.
- They compared $\rho_{i j}=0$ vs. $\rho_{i j}=\rho$.
- The problem is that standard asymptotics (normality and chi-sqare tests of fit) are not applicable as the number of parameters increases with the number of nodes.
- How to test $\rho_{i j}=\rho$ against a more complex model?


## Aside—The Normal Distribution was 275 Years Old on November 12, 2008!

Nos. 12. 1733-

## APPROXIMATIO AD

## Summam Terminorum binomil

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fere data fit, quod confideranti farile preebit, fequirur in Poecithet infinita quantiatem illam darum iri, eamque exhibiarumm nomerum illaun cujus Logarithmas hyperbolicus eft -1; hinc feet ut fif B detignarit numerum illum cujus Logarichmus byperbolies eft $-1+\frac{1}{12}-\frac{1}{300}+\frac{1}{1250}-\frac{1}{1050}$
 Seriei mutenour, ponaturque B xqualis numerocujus Logarithmus hyperbolicus eft $1-\frac{1}{12}+\frac{1}{j 50}-\frac{1}{1300}+\frac{1}{100_{0}}$ Sc. expreflio ill furura fit $\frac{1}{B \sqrt{3}}$.

Cum primum ad haso dilquifitionem animam appali, eo contentas fui ut prater propter valorem quastiutis B determinarem quod quidem fis Aum fierat additione paucorum bujas feriei Termisorum, quorum fumms fpeetaus foerat unguam Logarithmes iftios quantiatis, attamen uarditas convergentia me deterruerat quominu loogies procederem, donec vir DoAiffumus mihique aniciflamss yondas Strinay qui pot me ad hane difquifirionem methodo a mea valde diverfa fe costali, comperit quantiatem B denocare radicem quadratam circomierentiz Ciredi cejos ralus eft unitus, ita ut fi hacc circumétentia appelletur c, ratio Medii Termini ad Terninos omoes exprimetur per $\frac{1}{\sqrt{x}}$.
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$\overline{\pi+1-i \times} \log \cdot \overline{x+i-1}+\overline{\pi-1+i} \times \log \overline{n-1+2}=2 \log , m+\log \cdot \frac{n+1}{n}$

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## Hotelling and the Normal Distribution

Harold Hotelling, the "Normal Statistician", 1946


Display courtesy of Stephen Stigler.

## $p_{1}$ in Log-linear Form

- Probabilities for four situations between two nodes:
- Define

$$
y_{i j k l}= \begin{cases}1 & \text { if } D\left(x_{i j}, x_{j i}\right)=(k, l) \\ 0 & \text { otherwise }\end{cases}
$$

- Yields an $n \times n \times 2 \times 2$ 4-way array with "zeros" for $n \times n$ diagonals.
- Fienberg and Wasserman demonstrated that in the 4-way table, the log-linear model of no second-order interaction corresponds to $p_{1}$ with constant reciprocation, i.e., [12][13][14][23][24][34], and that the standard iterative proportional fitting algorithm.
- Edge-dependent reciprocation corresponds to log-linear model [12][134][234].


## Primer: Algebraic Representation and Computer Algebra Tools

- Log-linear models are represented by polynomial maps and parameter space is toric variety.
- Multinomial likelihood is a monomial.
- Fiber consists of all tables with margins $t$.
- Markov basis generates fiber and consists of minimal generators of toric ideal.
- Computer algebra tools:
- 4ti2
- We use 4 ti2 to generate basis elements (perhaps redundant) for Markov bases for specific values of $n$. Can use these to compute exact distribution given the MSSs.
- Polymake—Examines fiber for MLE existence.
- We use Polymake to explore conditions for MLEs to exist, as in example for $\rho_{i j}=0$. Affects computation and assessment of fit.


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$p_{i j}(a, b) \mapsto \lambda_{i j} \alpha_{i}^{a} \alpha_{j}^{b} \beta_{i}^{b} \beta_{j}^{a} \theta^{a+b} \rho_{i j}^{\min (a, b)}$ for all $i<j \in\{1, \ldots, n\}$


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- For example, $n=3$ and $\rho_{i j}$ edge-dependent, the variety is a degree-3 hypersurface in $\mathbb{P}^{11}$.
- Its defining ideal gives Markov basis,
- it will connect all networks with the same sufficient statistics.


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Theorem (FPR)
If $\rho_{i j}=0$, the ideal of the simplified model equals $I_{G_{n}}+T_{n}$ where $T_{n}$ is generated by $p_{i j}(1,0) p_{i j}(0,1)-p_{i j}(1,1)$ and $I_{G_{n}}$ is the toric ideal of the edge subring of $G_{n}:=K_{n, n} \backslash\{i, i\}$.

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## Theorem (FPR)

If $\rho_{i j}=\rho+\rho_{i}+\rho_{j}$, the ideal of the simplified model equals $I_{G_{n}}+Q_{n}$ where $I_{G_{n}}$ is as above, and $Q_{n}$ is the toric ideal of the edge subring of $K_{n}$.

## Main Theorem - toric ideal of $p_{1}$

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By multi-homogeneous, we mean with respect to each pair $i, j$.

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- In progress: Markov moves for all three cases of $\rho_{i j}$.


## Example with 4 nodes



Figure: A degree-5 binomial

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Figure: the corresponding path in $K_{4,4} \backslash\{i, i\}$

## 3-node network

For all 3 cases of $\rho_{i j}$, there is only one Markov move:

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$$

The corresponding Markov move is to remove edges $1 \rightarrow 2,2 \rightarrow 3$ and $3 \rightarrow 1$, and replace them by the same edges oriented in the opposite direction.

## 4-node network, $\rho_{i j}$ constant

- Reminder: $\rho_{i j}=\rho$, including $\rho_{i j}=0$.


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Figure: A binomial of degree 4

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represents

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$$

## 4-node network, $\rho_{i j}$ Constant



Figure: A more complicated binomial of degree 4
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$$

- The moves come in degrees $3,4,5,6$.


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Reminder: $\rho_{i j}=\rho+\rho_{i}+\rho_{j}$.

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What follows is a list of all possible moves on a 4-node network, with respect to symmetry of course.

## 4-node network, $\rho_{i j}$ edge-dependent



Figure: A binomial of degree 3

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Figure: A binomial of degree 4

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Figure: A binomial of degree 4

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## 4-node network, $\rho_{i j}$ edge-dependent



Figure: A binomial of degree 4

## 4-node network, $\rho_{i j}$ edge-dependent



Figure: A binomial of degree 4


Figure: A binomial of degree 5

## 4-node network, $\rho_{i j}$ edge-dependent



Figure: A binomial of degree 5

4-node network, $\rho_{i j}$ edge-dependent


Figure: A binomial of degree 5


Figure: A binomial of degree 6

## Summary

- Graphs and network models.
- $p_{1}$ and its algebraic representation:
- Reviewed of $p_{1}$ and log-linear models.
- Developed Markov bases.
- Markov bases and proposed uses:
- Existence of MLE.
- Generate exact distribution for $p_{1}$ and use for assessing goodness-of-fit.
- Future work includes:
- Completing algebraic characterization of $p_{1}$ models and putting them to use.
- Generalizations to exponential random graph models (ERGM) models [also known as $p^{*}$ models]:
- Many complex statistical issues including existence of MLEs (non-existence $=$ degeneracies involving zero estimates) and near-degeneracies.
- Scaling tools up to be of use for analysis large networks with efficient computation.
... The End ...

