

# ADDITIVE AND MULTIPLICATIVE LATENT FACTOR MODELS FOR NETWORK INFERENCE

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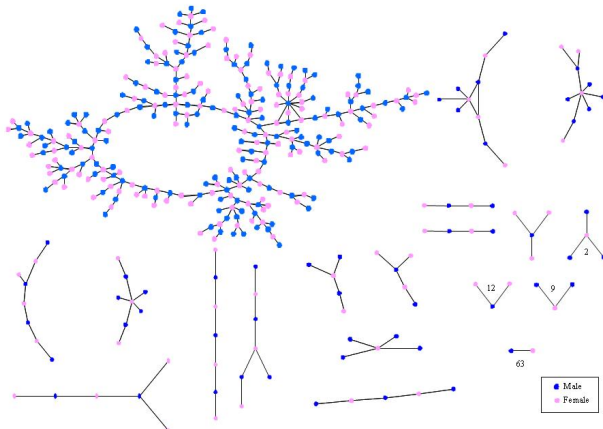
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# Networks are important; How to Study

Obligatory Network Graph Here. This is Sexual Network in Typical Midwest High School (Bearman, Moody, Stovel, 2004).



# Promise of Exponential Random Graph Models

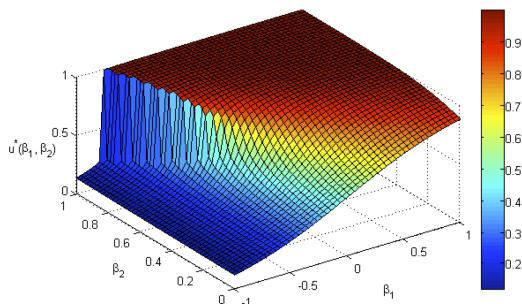
- Early 1970s development of pseudolikelihood estimation: Ove Frank (1971); Julian Besag (1972) proposed using a logistic regression with network characteristics as covariates. Birth of ERGM.
- $P(x) = \exp(\Theta^T s(x) - \psi(\theta))$ , where  $x$  is an adjacency matrix that is a graph,  $s(\theta)$  are some set of sufficient statistics for the graph, and  $\psi(\theta)$  is a normalizing constant, often set to be  $\log \sum_x e^{\theta^T s(x)}$ . This is often estimated via pseudolikelihood, simply by regressing  $x \sim \text{logit}(z(x))$ .
- Maximum Likelihood is a better approach with Robbins and Monro, or the importance sampling approach of Geyer & Thompson. More advances with Bayesian approaches are available now with MCMC (Koskinen, Robins & Pattison).
- In the 1990s, networks became more widely recognized as important and the ERGM approach was often employed to estimate models in a variety of network domains. Needle sharing communities, HIV infections, for example.

# There is a problem with ERGM.

- Schweinberger, M. (2011). Instability, sensitivity, and degeneracy of discrete exponential families. **Journal of the American Statistical Association**, 106(496):1361–1370.
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- Chatterjee, S. and Diaconis, P. (2013). Estimating and understanding exponential random graph models. **The Annals of Statistics**, 41(5):2428–2461.
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# It is a bug, not a feature.

- Probabilistic ERGM models place almost all of the probability on networks that are either nearly empty (degenerate) with no linkages or nearly saturated with all nodes being interconnected.
- The likelihood surface contains steep or discontinuous gradients that render it impossible to solve numerically (or analytically). Even (especially) for very small networks this is problematic.



# Let's (re)start with the data and build up an approach

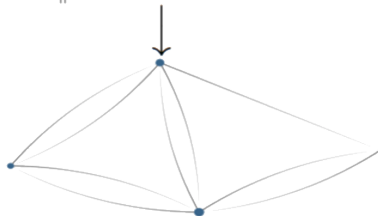
Relational data consists of

- a set of units or nodes
- a set of measurements,  $y_{ij}$ , specific to pairs of nodes  $(i, j)$

Sender	Receiver	Event
$i$	$j$	$y_{ij}$
$\vdots$	$k$	$y_{ik}$
	$l$	$y_{il}$
$j$	$i$	$y_{ji}$
$\vdots$	$k$	$y_{jk}$
	$l$	$y_{jl}$
$k$	$i$	$y_{ki}$
$\vdots$	$j$	$y_{kj}$
	$l$	$y_{kl}$
$l$	$i$	$y_{li}$
$\vdots$	$j$	$y_{lj}$
	$k$	$y_{lk}$



	$i$	$j$	$k$	$l$
$i$	NA	$y_{ij}$	$y_{ik}$	$y_{il}$
$j$	$y_{ji}$	NA	$y_{jk}$	$y_{jl}$
$k$	$y_{ki}$	$y_{kj}$	NA	$y_{kl}$
$l$	$y_{li}$	$y_{lj}$	$y_{lk}$	NA



# Relational data assumptions

GLM:  $y_{ij} \sim \beta^T X_{ij} + e_{ij}$

Networks typically show evidence against independence of  $e_{ij} : i \neq j$

Not accounting for dependence can lead to:

- biased effects estimation
- uncalibrated confidence intervals
- poor predictive performance
- inaccurate description of network phenomena

We've been hearing this concern for decades now:

Thompson & Walker (1982)	Beck et al. (1998)	Franzese & Hays (2007)
Frank & Strauss (1986)	Signorino (1999)	Snijders (2011)
Kenny (1996)	Li & Loken (2002)	Aronow et al. (2015)
Krackhardt (1998)	Hoff & Ward (2004)	Athey et al. (2016)

- Nodal and dyadic dependencies in networks
  - Can model using the “A” in AME
- Third order dependencies
  - Can model using the “M” in AME
- Application



# What network phenomena? Sender heterogeneity

Values across a row, say  $\{y_{ij}, y_{ik}, y_{il}\}$ , may be more similar to each other than other values in the adjacency matrix because each of these values has a common sender  $i$

	$i$	$j$	$k$	$l$
$i$	NA	$y_{ij}$	$y_{ik}$	$y_{il}$
$j$	$y_{ji}$	NA	$y_{jk}$	$y_{jl}$
$k$	$y_{ki}$	$y_{kj}$	NA	$y_{kl}$
$l$	$y_{li}$	$y_{lj}$	$y_{lk}$	NA

# What network phenomena? Receiver heterogeneity

Values across a column, say  $\{y_{ji}, y_{ki}, y_{li}\}$ , may be more similar to each other than other values in the adjacency matrix because each of these values has a common receiver  $i$

	$i$	$j$	$k$	$l$
$i$	NA	$y_{ij}$	$y_{ik}$	$y_{il}$
$j$	$y_{ji}$	NA	$y_{jk}$	$y_{jl}$
$k$	$y_{ki}$	$y_{kj}$	NA	$y_{kl}$
$l$	$y_{li}$	$y_{lj}$	$y_{lk}$	NA

# What network phenomena? Sender-Receiver Covariance

Actors who are more likely to send ties in a network may also be more likely to receive them

	$i$	$j$	$k$	$l$
$i$	NA	$y_{ij}$	$y_{ik}$	$y_{il}$
$j$	$y_{ji}$	NA	$y_{jk}$	$y_{jl}$
$k$	$y_{ki}$	$y_{kj}$	NA	$y_{kl}$
$l$	$y_{li}$	$y_{lj}$	$y_{lk}$	NA

# What network phenomena? Reciprocity

Values of  $y_{ij}$  and  $y_{ji}$  may be statistically dependent

	$i$	$j$	$k$	$l$
$i$	NA	$y_{ij}$	$y_{ik}$	$y_{il}$
$j$	$y_{ji}$	NA	$y_{jk}$	$y_{jl}$
$k$	$y_{ki}$	$y_{kj}$	NA	$y_{kl}$
$l$	$y_{li}$	$y_{lj}$	$y_{lk}$	NA

# Social Relations Model (The “A” in AME)

We use this model to form the additive effects portion of AME

$$y_{ij} = \mu + e_{ij}$$

$$e_{ij} = a_i + b_j + \epsilon_{ij}$$

$$\{(a_1, b_1), \dots, (a_n, b_n)\} \sim N(0, \Sigma_{ab})$$

$$\{(\epsilon_{ij}, \epsilon_{ji}) : i \neq j\} \sim N(0, \Sigma_{\epsilon}), \text{ where}$$

$$\Sigma_{ab} = \begin{pmatrix} \sigma_a^2 & \sigma_{ab} \\ \sigma_{ab} & \sigma_b^2 \end{pmatrix} \quad \Sigma_{\epsilon} = \sigma_{\epsilon}^2 \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix}$$

- $\mu$  baseline measure of network activity
- $e_{ij}$  residual variation that we will use the SRM to decompose

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- row/sender effect ( $a_i$ ) & column/receiver effect ( $b_j$ )
- Modeled jointly to account for correlation in how active an actor is in sending and receiving ties

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$$\{(a_1, b_1), \dots, (a_n, b_n)\} \sim N(0, \Sigma_{ab})$$

$$\{(\epsilon_{ij}, \epsilon_{ji}) : i \neq j\} \sim N(0, \Sigma_\epsilon), \text{ where}$$

$$\Sigma_{ab} = \begin{pmatrix} \sigma_a^2 & \sigma_{ab} \\ \sigma_{ab} & \sigma_b^2 \end{pmatrix} \quad \Sigma_\epsilon = \sigma_\epsilon^2 \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix}$$

- $\sigma_a^2$  and  $\sigma_b^2$  capture heterogeneity in the row and column means
- $\sigma_{ab}$  describes the linear relationship between these two effects (i.e., whether actors who send [receive] a lot of ties also receive [send] a lot of ties)

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$$y_{ij} = \mu + e_{ij}$$

$$e_{ij} = a_i + b_j + \epsilon_{ij}$$

$$\{(a_1, b_1), \dots, (a_n, b_n)\} \sim N(0, \Sigma_{ab})$$

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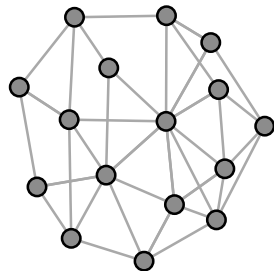
$$\Sigma_{ab} = \begin{pmatrix} \sigma_a^2 & \sigma_{ab} \\ \sigma_{ab} & \sigma_b^2 \end{pmatrix} \quad \Sigma_{\epsilon} = \sigma_{\epsilon}^2 \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix}$$

- $\epsilon_{ij}$  captures the within dyad effect
- Second-order dependencies are described by  $\sigma_{\epsilon}^2$
- Reciprocity, aka within dyad correlation, represented by  $\rho$

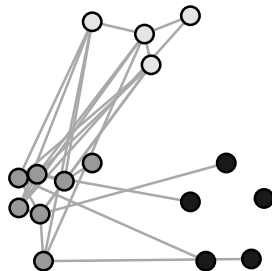


# Third Order Dependencies

HOMOPHILY



STOCHASTIC EQUIVALENCE



To account for these patterns we can build on what we have so far and find an expression for  $\gamma$ :

$$y_{ij} \approx \beta^T X_{ij} + a_i + b_j + \gamma(u_i, v_j)$$

# Latent Factor Model: The “M” in AME

Each node  $i$  has an unknown latent factor

$$\mathbf{u}_i, \mathbf{v}_j \in \mathbb{R}^k \quad i, j \in \{1, \dots, n\}$$

The probability of a tie from  $i$  to  $j$  depends on their latent factors

$$\begin{aligned}\gamma(\mathbf{u}_i, \mathbf{v}_j) &= \mathbf{u}_i^T D \mathbf{v}_j \\ &= \sum_{k \in K} d_k u_{ik} v_{jk}\end{aligned}$$

$D$  is a  $K \times K$  diagonal matrix

Can account for both stochastic equivalence and homophily.

# Inner Product versus Euclidean Distance

We focus on two approaches to the latent space: the latent distance model (LDM) and the latent factor model (LFM).

Latent distance model

$$\alpha(\mathbf{u}_i, \mathbf{u}_j) = -|\mathbf{u}_i - \mathbf{u}_j|$$
$$\mathbf{u}_i \in \mathbb{R}^K, \quad i \in \{1, \dots, n\}$$

Latent factor model

$$\alpha(\mathbf{u}_i, \mathbf{u}_j) = \mathbf{u}_i^\top \Lambda \mathbf{u}_j$$
$$\mathbf{u}_i \in \mathbb{R}^K, \quad i \in \{1, \dots, n\}$$

$\Lambda$  a  $K \times K$  diagonal matrix

(1)

# Putting it all together

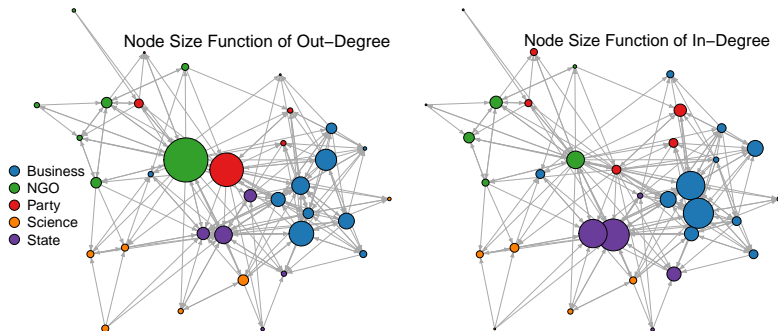
The AME approach can be restated as simple (*simple*) regression

$$\begin{aligned}y_{ij} &= g(\theta_{ij}) \\ \theta_{ij} &= \boldsymbol{\beta}^\top \mathbf{X}_{ij} + e_{ij} \\ e_{ij} &= a_i + b_j + \epsilon_{ij} + \alpha(\mathbf{u}_i, \mathbf{v}_j) , \text{ where} \\ \alpha(\mathbf{u}_i, \mathbf{v}_j) &= \mathbf{u}_i^\top \mathbf{D} \mathbf{v}_j = \sum_{k \in K} d_k u_{ik} v_{jk}.\end{aligned}\tag{2}$$

The `amen` package implements this. Let's use it to hit some nails.

# Swiss Climate Change Application

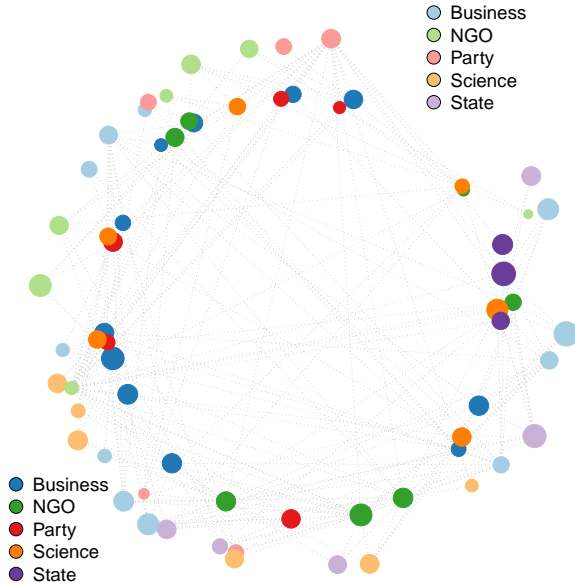
Cross-sectional network measuring whether an actor indicated that they collaborated with another during the policy design of the Swiss CO<sub>2</sub> act (Ingold 2008)



# Parameter Estimates

	Expected Effect	Logit	MRQAP	LDM	ERGM	AME
<b>Conflicting policy preferences</b>						
Business vs. NGO	—	-0.86	-0.87*	-1.37*	-1.11*	-1.37*
Opposition/alliance	+	1.21*	1.14*	0.00	1.22*	1.08*
Preference dissimilarity	—	-0.07	-0.60	-1.76*	-0.44	-0.79*
<b>Transaction costs</b>						
Joint forum participation	+	0.88*	0.75*	1.51*	0.90*	0.92*
<b>Influence</b>						
Influence attribution	+	1.20*	1.29*	0.08	1.00*	1.09*
Alter's influence indegree	+	0.10*	0.11*	0.01	0.21*	0.11*
Influence absolute diff.	—	-0.03*	-0.06*	0.04	-0.05*	-0.07*
Alter = Government actor	+	0.63*	0.68	-0.46	1.04*	0.55
<b>Functional requirements</b>						
Ego = Environmental NGO	+	0.88*	0.99	-0.60	0.79*	0.67
Same actor type	+	0.74*	1.12*	1.17*	0.99*	1.04*

# Latent Factor Visualization



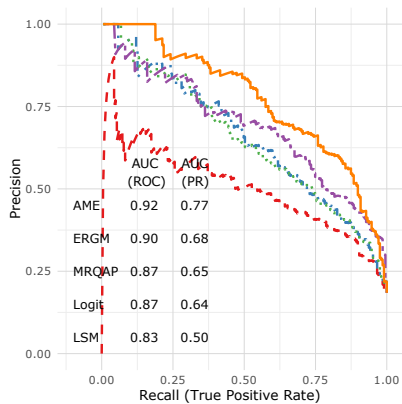
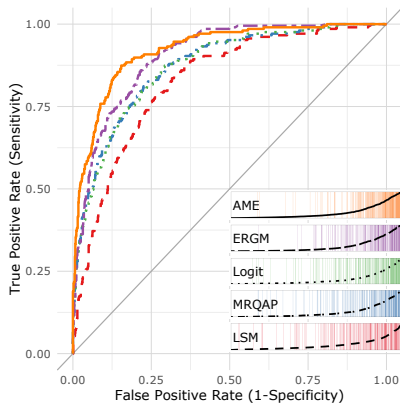
# Out of Sample Performance Assessment

- Randomly divide the  $n \times (n - 1)$  data points into  $S$  sets of roughly equal size, letting  $s_{ij}$  be the set to which pair  $\{ij\}$  is assigned.
- For each  $s \in \{1, \dots, S\}$ :
  - Obtain estimates of the model parameters conditional on  $\{y_{ij} : s_{ij} \neq s\}$ , the data on pairs not in set  $s$ .
  - For pairs  $\{kl\}$  in set  $s$ , let  $\hat{y}_{kl} = E[y_{kl} | \{y_{ij} : s_{ij} \neq s\}]$ , the predicted value of  $y_{kl}$  obtained using data not in set  $s$ .

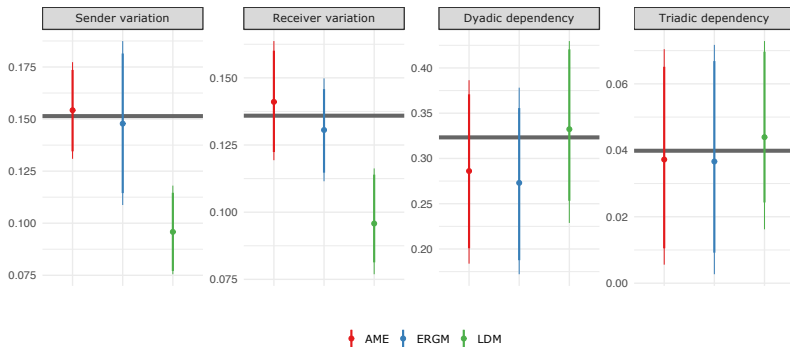
This procedure generates a sociomatrix of out-of-sample predictions of the observed data



# Performance Comparison



# Network Dependencies



# Conclusion

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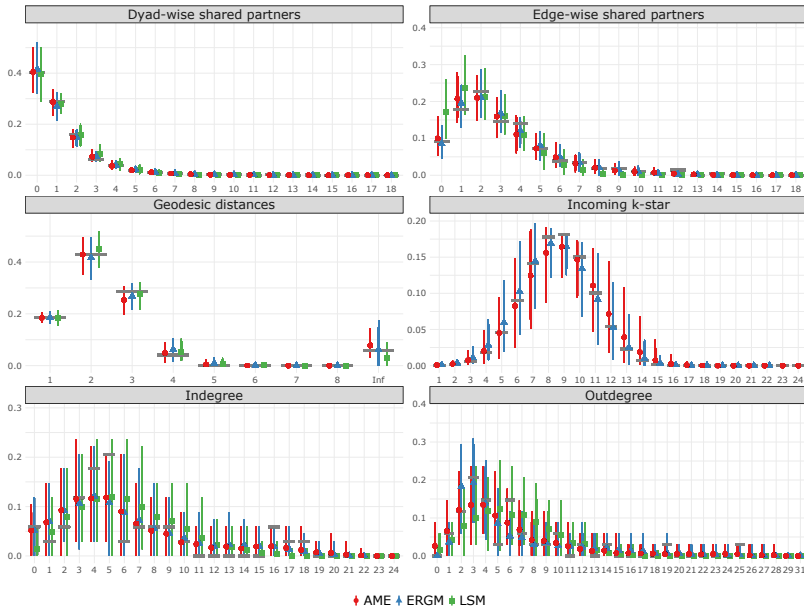
1. AME works for binary, count, and continuous relational data.
2. AME allows longitudinal network data (i.e., tensors), wherein there can be different observations (nodes) in different time slices.
3. AME is a regression based approach that has a numerically tractable likelihood function and it is not threatened by missing data.
4. Is available on CRAN, with some of the more recent features to be delivered later this summer, but available to interested beta testers.

# References

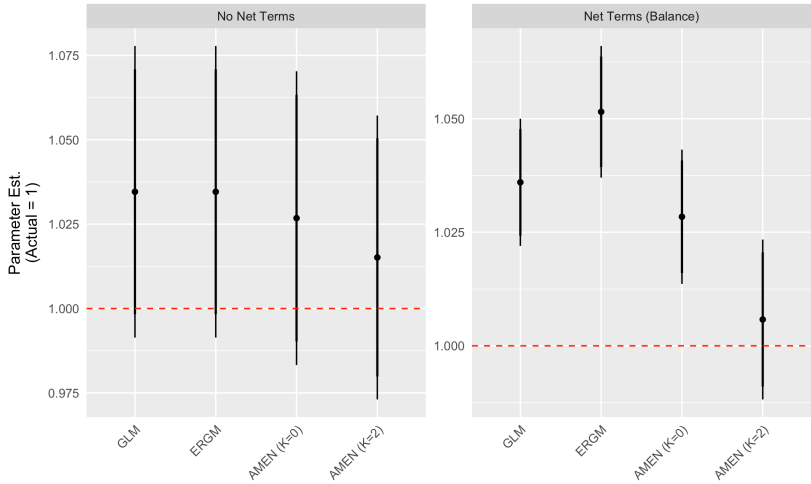
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THANKS.

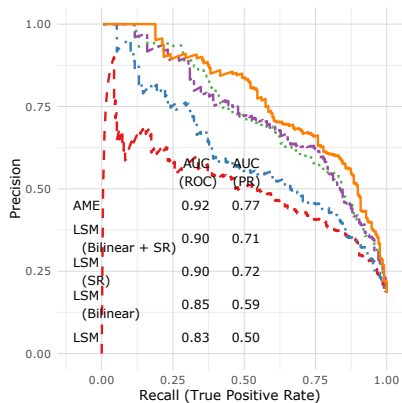
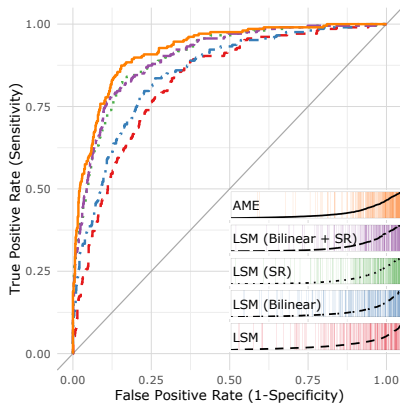
# Standard Network Dependence Measures



# Simulation Comparison

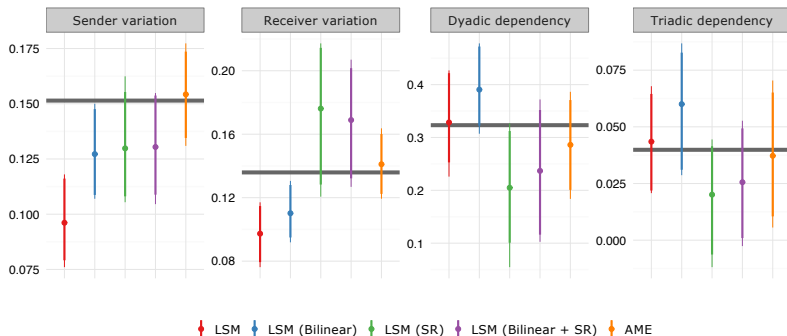


# AMEN v LSM Performance

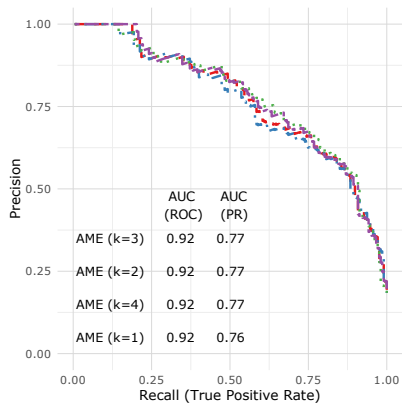
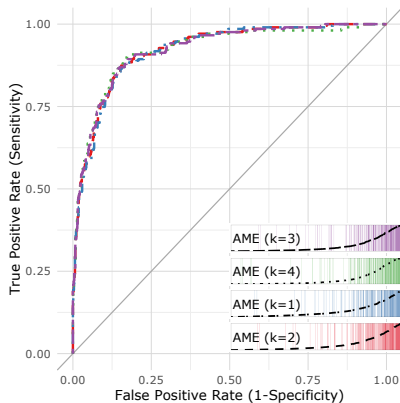




# AMEN versus LSM Net Dependence



# AMEN varying $K$ Performance



# AMEN varying $K$ Net Dependence

