

# Network analysis with the stochastic block model (using variational approximations)

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# Outline

- 1 State-space models for networks (inc. SBM)
- 2 Variational inference (inc. SBM)
- 3 Extensions of SBM
- 4 (Variational) Bayesian model averaging
- 5 Towards  $W$ -graphs
- 6 Goodness-of-fit using network motif

# Heterogeneity in interaction networks

# Understanding network structure

Networks describe interactions between entities.

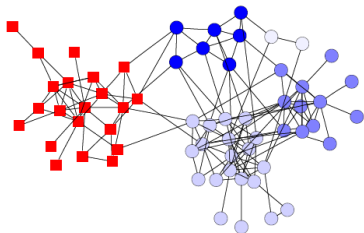
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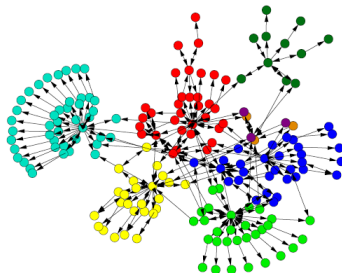
Observed networks display heterogeneous topologies, that one would like to decipher and better understand.

Dolphine social network.



[Newman and Girvan (2004)]

Hyperlink network.



# Modeling network heterogeneity

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**General setting for binary graphs.** [Bollobás *et al.* (2007)]:

- A latent (unobserved) variable  $Z_i$  is associated with each node:

$$\{Z_i\} \text{ iid } \sim \pi$$

- Edges  $Y_{ij} = \mathbb{I}\{i \sim j\}$  are independent conditionally to the  $Z_i$ 's:

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**We focus here on model approaches**, in contrast with, e.g.

- Graph clustering [Girvan and Newman (2002)], [Newman (2004)];
- Spectral clustering [von Luxburg *et al.* (2008)].



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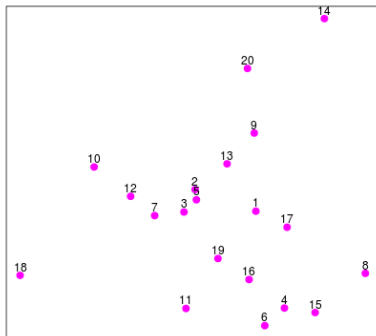
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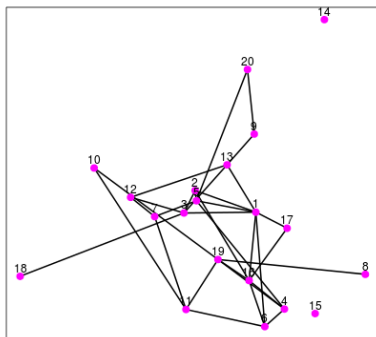
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$$Y = \begin{pmatrix} 0 & 1 & 1 & 0 & 1 & \dots \\ 0 & 0 & 1 & 0 & 1 & \dots \\ 0 & 0 & 0 & 0 & 0 & \dots \\ 0 & 0 & 0 & 0 & 1 & \dots \\ 0 & 0 & 0 & 0 & 0 & \dots \\ \vdots & \vdots & \vdots & \vdots & \vdots & \ddots \end{pmatrix}$$

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# A variety of state-space models

## Continuous. Latent position models.

- [Hoff *et al.* (2002)]:

$$Z_i \in \mathbb{R}^d, \quad \text{logit}[\gamma(z, z')] = a - |z - z'|$$

- [Handcock *et al.* (2007)]:

$$Z_i \sim \sum_k p_k \mathcal{N}_d(\mu_k, \sigma_k^2 I)$$

- [Lovász and Szegedy (2006)]:

$$Z_i \sim \mathcal{U}_{[0,1]}, \quad \gamma(z, z') : [0, 1]^2 \rightarrow [0, 1] =: \text{graphon function}$$

- [Daudin *et al.* (2010)]:

$$Z_i \in \mathcal{S}_K, \quad \gamma(z, z') = \sum_{k,\ell} z_k z'_\ell \gamma_{k\ell}$$

# Stochastic Block Model (SBM)

A mixture model for random graphs.

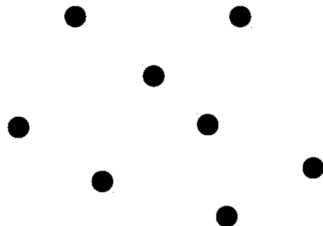
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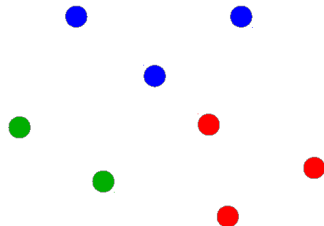
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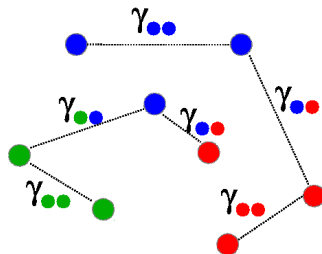
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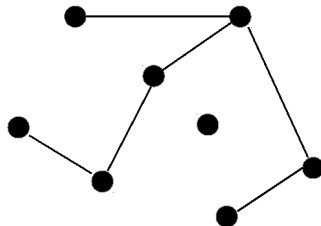
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# Variational inference

# Incomplete data models

**Aim.** Based on the observed network  $Y = (Y_{ij})$ , we want to infer

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State space models belong to the class of incomplete data models as

- the edges  $(Y_{ij})$  are observed,
- the latent positions (or status)  $(Z_i)$  are not.

→ usual issue in unsupervised classification.

# Frequentist maximum likelihood inference

**Likelihood.** The (log-)likelihood

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where  $\mathcal{H}$  stands for the entropy.

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Aims at maximizing the log-likelihood

$$\log P(Y; \theta)$$

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→ sometimes impossible (SBM: ...)



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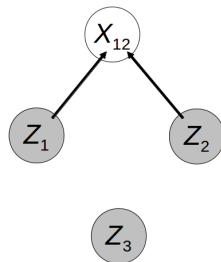
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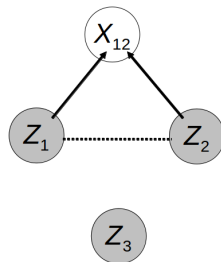
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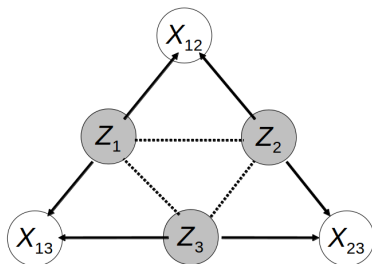


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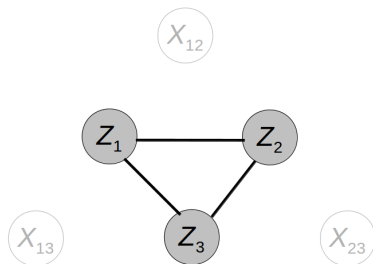


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**Conditional distribution.** The dependency graph of  $Z$  given  $Y$  is a clique.

- No factorization can be hoped (unlike for HMM).
- $P(Z|Y; \theta)$  can not be computed (efficiently).
- Variational techniques may help as they provide

$$Q(Z) \simeq P(Z|Y).$$

# Variational inference

**Lower bound of the log-likelihood.** For any distribution  $Q(Z)$  [Jaakkola (2000), Wainwright and Jordan (2008)],

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**Link with EM.** This is similar to

$$\log P(Y) = \mathbb{E}[\log P(Y, Z)|Y] + \mathcal{H}[P(Z|Y)]$$

replacing  $P(Z|Y)$  with  $Q(Z)$ .

# Variational EM algorithm

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→ Taking  $\mathcal{Q} = \{\text{all possible distributions}\}$  gives  $Q^*(Z) = P(Z|Y)$  ... like EM does.

→ Variational approximations rely on the choice a set  $\mathcal{Q}$  of 'good' and 'tractable' distributions.

# Variational EM for SBM [Daudin *et al.* (2008)]

**Distribution class.**  $\mathcal{Q}$  = set of factorisable distributions:

$$\mathcal{Q} = \{Q : Q(Z) = \prod_i Q_i(Z_i)\}, \quad Q_i(Z_i) = \prod_k \tau_{ik}^{Z_{ik}}.$$

→ The approximate joint distribution is  $Q(Z_i, Z_j) = Q_i(Z_i)Q_j(Z_j)$ .

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The optimal approximation within this class satisfies a fix-point relation:

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also known as mean-field approximation in physics [Parisi (1988)].

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## Variational estimates.

- No general statistical guaranty for variational estimates.
- SBM is a very specific case for which the variational approximation is asymptotically exact [Celisse *et al.* (2012), Mariadassou and Matias (2015)].



# Variational Bayes inference

Bayesian perspective.

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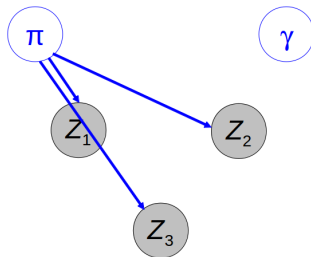
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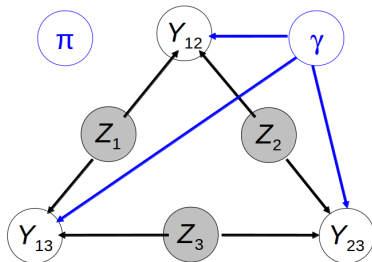
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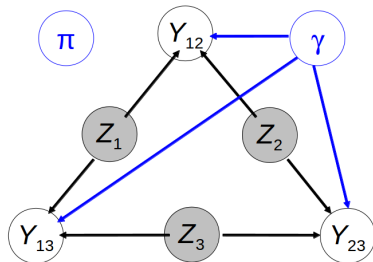
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Bayesian inference. The aim is then to get the joint conditional distribution of the parameters and of the hidden variables:

$$P(\theta, Z|Y).$$

# Variational Bayes algorithm

**Variational Bayes.** As  $P(\theta, Z|Y)$  is intractable, one look formula

$$Q^*(\theta, Z) = \arg \min_{Q \in \mathcal{Q}} KL [Q(\theta, Z) || P(\theta, Z|Y)]$$

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**Variational Bayes EM (VBEM).** When

- $P(Z, Y|\theta)$  belongs to the exponential family,
- $P(\theta)$  is the corresponding conjugate prior,

$Q^*$  can be obtained iteratively as [Beal and Ghahramani (2003)]

$$\log Q_\theta^h(\theta) \propto \mathbb{E}_{Q_Z^{h-1}} [\log P(Z, Y, \theta)], \quad \log Q_Z^h(Z) \propto \mathbb{E}_{Q_\theta^h} [\log P(Z, Y, \theta)].$$

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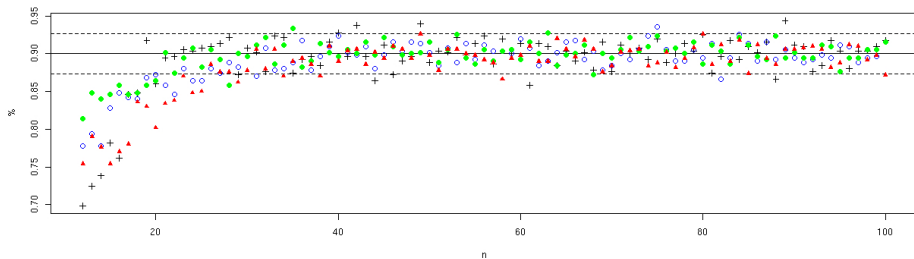
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Application to SBM: [Latouche *et al.* (2012)]



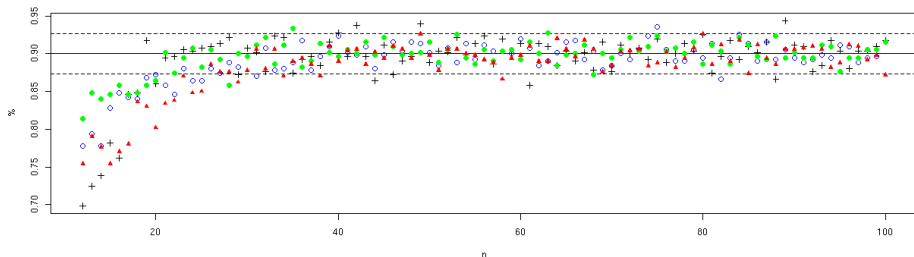
# VBEM: Simulation study [Gazal *et al.* (2012)]

Credibility intervals:  $\pi_1$ : +,  $\gamma_{11}$ :  $\triangle$ ,  $\gamma_{12}$ :  $\circ$ ,  $\gamma_{22}$ :  $\bullet$

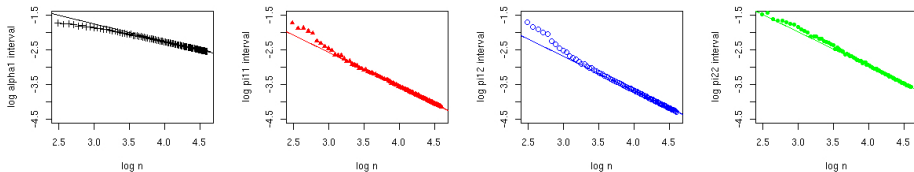


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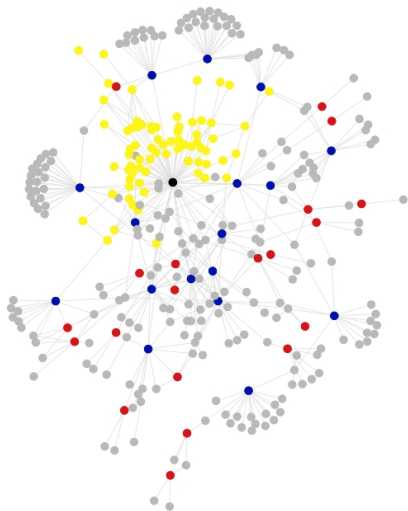
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Width of the posterior credibility intervals.  $\pi_1$ ,  $\gamma_{11}$ ,  $\gamma_{12}$ ,  $\gamma_{22}$

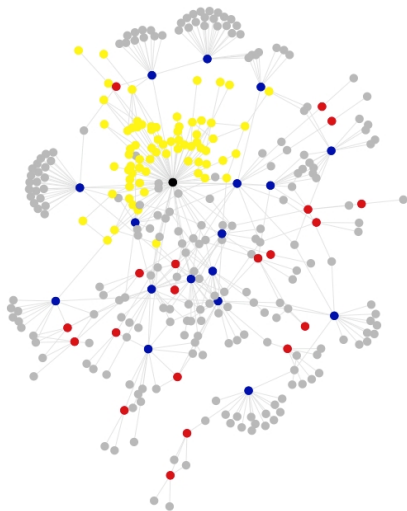


# SBM analysis of *E. coli* operon networks

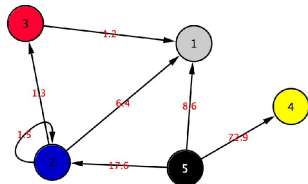


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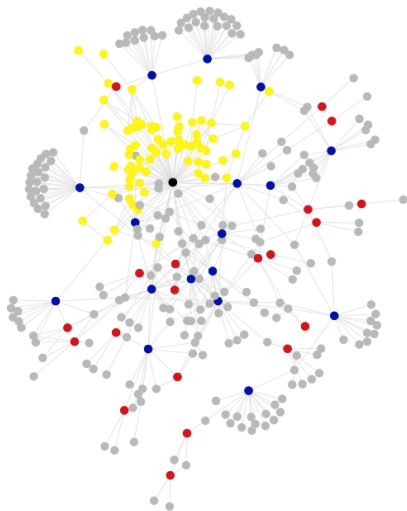


Meta-graph representation.



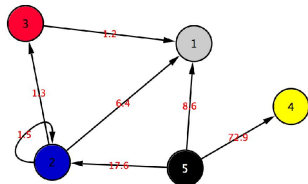
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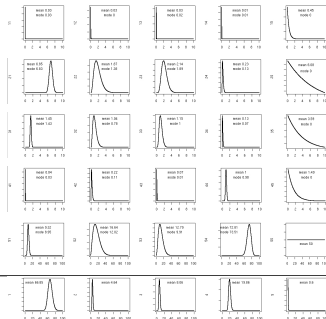


[Picard *et al.* (2009)]

Meta-graph representation.



Parameter estimates.  $K = 5$



# Some extensions of SBM

# GLM framework [Mariadassou *et al.* (2010)]

SBM can be combined with generalized linear models (GLM) to deal with both valued graphs and covariates.

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**Valued graphs.** Simply adapt the emission distribution:

$$(Y_{ij} \mid Z_i = k, Z_j = \ell) \sim F_{k\ell}(\cdot) := F(\cdot; \gamma_{k\ell})$$

where  $F$  = some (parametric) distribution: Bernoulli (regular SBM), Poisson, Gaussian, etc.



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**Covariates.** In the context of exponential family, covariates  $\mathbf{x}$  can be accounted for via a regression term

$$g(\mathbb{E}Y_{ij} \mid Z_i = k, Z_j = \ell) = \gamma_{k\ell} + \mathbf{x}_{ij}\beta$$

where

- $g$  stands for the link function (logit, log, identity, etc.);
- $\beta$  may depend or not on the groups ( $\beta \rightarrow \beta_{k\ell}$ ).

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**Data:**  $n = 51$  tree species,  
 $X_{ij}$  = number of shared parasites  
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$e^{\hat{\gamma}_{k\ell}}$	T1	T2	T3	T4	T5	T6	T7
T1	14.46	4.19	5.99	7.67	2.44	0.13	1.43
T2		14.13	0.68	2.79	4.84	0.53	1.54
T3			3.19	4.10	0.66	0.02	0.69
T4				7.42	2.57	0.04	1.05
T5					3.64	0.23	0.83
T6						0.04	0.06
T7							0.27
$\hat{\pi}_k$	7.8	7.8	13.7	13.7	15.7	19.6	21.6

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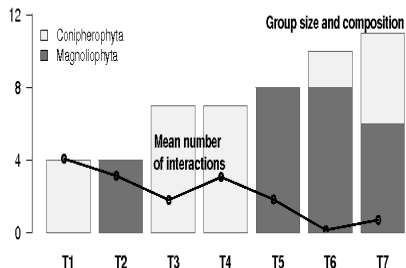
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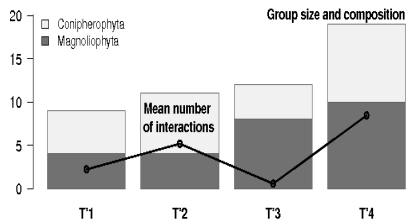
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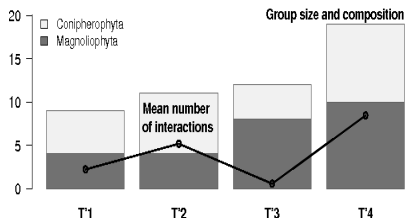
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→ Groups are no longer associated with the phylogenetic structure.

→ Mixture = residual heterogeneity of the regression.

# (Variational) Bayesian model averaging

# Model choice

**Model selection.** The number of classes  $K$  generally needs to be estimated.

- In the frequentist setting, an approximate ICL criterion can be derived

$$ICL = \mathbb{E}[\log P(Y, Z)|Y] - \frac{1}{2} \left\{ \frac{K(K+1)}{2} \log \frac{n(n-1)}{2} - (K-1) \log n \right\}.$$

- In the Bayesian setting, exact versions of BIC and ICL criteria can be calculated as

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**But**, in some applications, it may be useless or meaningless and model averaging may be preferred.

# Bayesian model averaging (BMA)

General principle. [Hoeting *et al.* (1999)]

- $\Delta$ : a parameter that can be defined under a series of different models  $\{\mathcal{M}_K\}_K$ .
- Denote  $P_K(\Delta|Y)$  its posterior distribution under model  $\mathcal{M}_K$ .

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**Remarks.**

- $w_K = P(\mathcal{M}_K|Y)$ : weight given to model  $\mathcal{M}_K$  for the estimation of  $\Delta$ .
- Calculating of  $w_K$  is not easy, but variational approximation may help.



# Variational Bayesian model averaging [Volant *et al.* (2012)]

## Variational Bayes formulation.

- $\mathcal{M}_K$  can be viewed as one more hidden layer
- Variational Bayes then aims at finding

$$Q^*(K, \theta, Z) = \arg \min_{Q \in \mathcal{Q}} KL[Q(K, \theta, Z) || P(K, \theta, Z | Y)]$$

with  $\mathcal{Q} = \{Q(\theta, Z) = Q_\theta(\theta | K) Q_Z(Z | K) Q_K(K)\}^1$

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## Optimal variational weights:

$$\begin{aligned} Q_K^*(K) &\propto P(K) \exp\{\log P(Y|K) - KL[Q^*(Z, \theta | K); P(Z, \theta | Y, K)]\} \\ &= P(K|Y) e^{-KL[Q^*(Z, \theta | K); P(Z, \theta | Y, K)]}. \end{aligned}$$

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# VBMA: the recipe

For  $K = 1 \dots K_{\max}$

- Use regular VBEM to compute the approximate conditional posterior

$$Q_{Z,\theta|K}^*(Z, \theta|K) = Q_{Z|K}^*(Z|K)Q_{\theta|K}^*(\theta|K);$$

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Deduce the approximate posterior of  $\Delta$ :

$$P(\Delta|Y) \approx \sum_K w_K Q_{\theta|K}^*(\Delta|K)$$

# Towards $W$ -graphs

# $W$ -graph model

Graphon function  $\gamma(z, z')$

$W$  graph.

Latent variables:

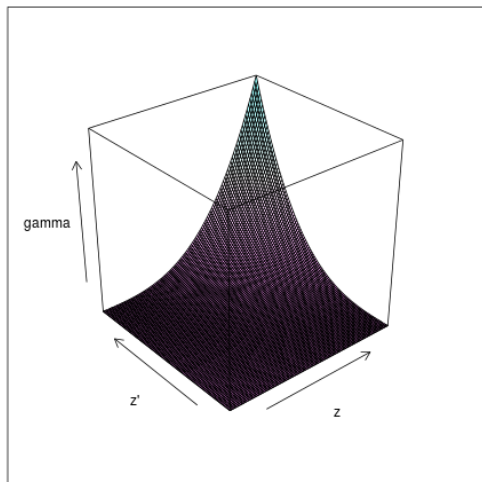
$$(Z_i) \text{ iid } \sim \mathcal{U}_{[0,1]},$$

Graphon function  $\gamma$ :

$$\gamma(z, z') : [0, 1]^2 \rightarrow [0, 1]$$

Edges:

$$\Pr\{Y_{ij} = 1\} = \gamma(Z_i, Z_j)$$



# Inference of the graphon function

## Probabilistic point of view.

- $W$ -graph have been mostly studied in the probability literature: [Lovász and Szegedy (2006)], [Diaconis and Janson (2008)]
- Motif (sub-graph) frequencies are invariant characteristics of a  $W$ -graph.
- Intrinsic un-identifiability of the graphon function  $\gamma$  is often overcome by imposing that  $u \mapsto \int \gamma(u, v) dv$  is monotonous increasing.



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## Statistical point of view.

- Not much attention has been paid to its inference until very recently: [Chatterjee (2012)], [Airoldi *et al.* (2013)], ...
- The latter also uses SBM as a proxy for  $W$ -graph.

# SBM as a $W$ -graph model

Latent variables:

$$(U_i) \text{ iid } \sim \mathcal{U}[0, 1]$$

$$Z_{ik} = \mathbb{I}\{\sigma_{k-1} \leq U_i < \sigma_k\}$$

where  $\sigma_k = \sum_{\ell=1}^k \pi_\ell$ .

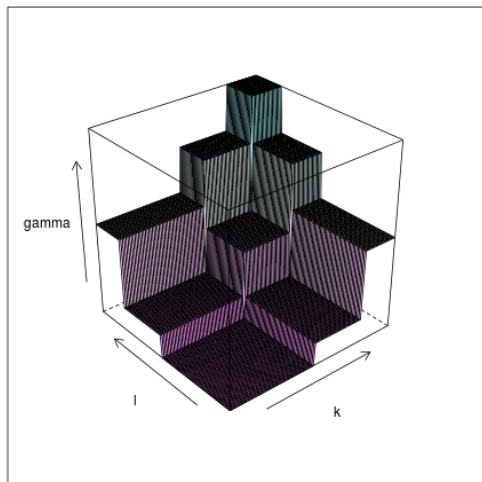
Blockwise constant graphon:

$$\gamma(z, z') = \gamma_{kl}$$

Edges:

$$\Pr\{Y_{ij} = 1\} = \gamma(Z_i, Z_j)$$

Graphon function  $\gamma_K^{SBM}(z, z')$



# Variational Bayes estimation of $\gamma(z, z')$ [Latouche and R. (2013)]

Posterior mean of  $\gamma_K^{SBM}(z, z')$

**VBEM inference** provides the approximate posteriors:

$$(\pi|Y) \approx \text{Dir}(\pi^*)$$

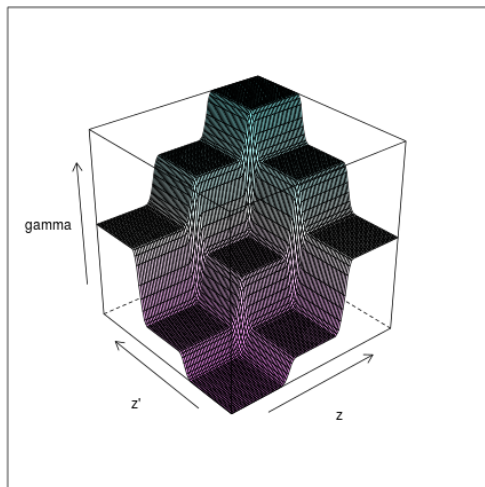
$$(\gamma_{kl}|Y) \approx \text{Beta}(\gamma_{kl}^{0*}, \gamma_{kl}^{1*})$$

**Estimate of  $\gamma(u, v)$ .**

$$\hat{\gamma}_K^{SBM}(u, v) = \tilde{\mathbb{E}}(\gamma_{C(u), C(v)} | Y)$$

where  $C(u) = 1 + \sum_k \mathbb{I}\{\sigma_k \leq u\}$ .

[Gouda and Szántai (2010)]



# Model averaging

**Model averaging:** There is no 'true  $K$ ' in the  $W$ -graph model.

**Apply VBMA recipe.** For  $K = 1..K_{\max}$ , fit an SBM model via VBEM and compute

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Perform model averaging as

$$\hat{\gamma}(z, z') = \sum_K w_K \hat{\gamma}_K^{SBM}(z, z')$$

where  $w_K$  is the variational weights arising from variational Bayes inference.

# Some simulations

**Design.** Symetric graphon:

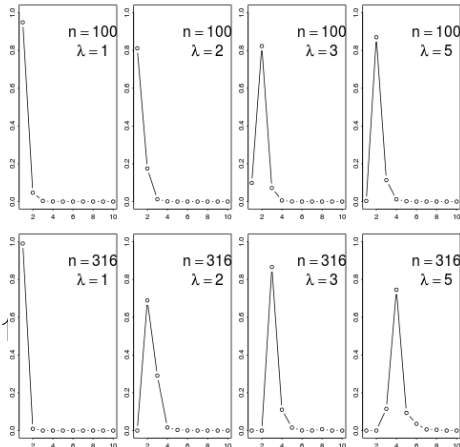
$$\gamma(u, v) = \rho \lambda^2 (uv)^{\lambda-1}$$

- $\lambda \uparrow$ : imbalanced graph
- $\rho \uparrow$ : dense graph

**Results.**

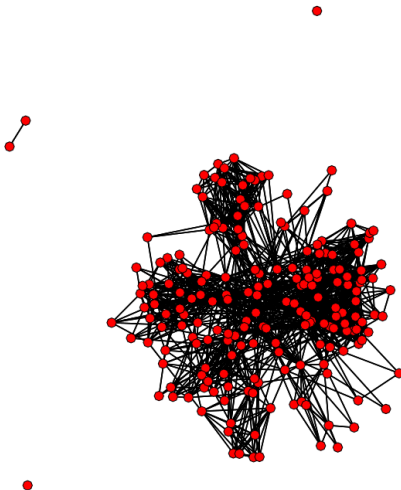
- More complex models as  $n$  and  $\lambda$   $\uparrow$
- Posterior fairly concentrated

Variational posterior for  $K$ :  $Q^*(K)$ .



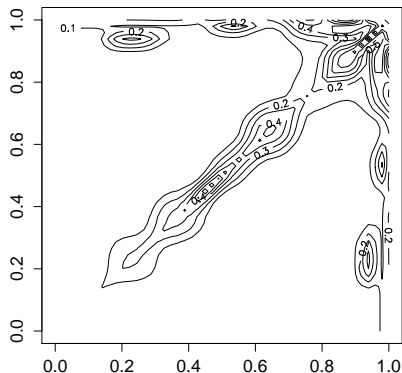
# French political blogosphere

Website network. French political blogs: 196 nodes, 1432 edges.



# French political blogosphere

Infered graphon.  $\widehat{W}(u, v) = \mathbb{E}(\gamma(u, v) | Y)$

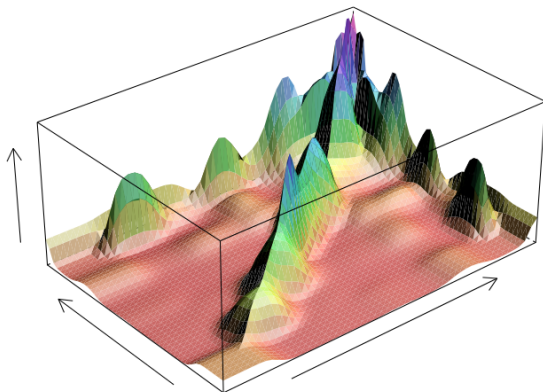


Motif probability can be estimated as well as  $\widehat{\mu}(m) = \mathbb{E}(\mu(m) | Y)$ .



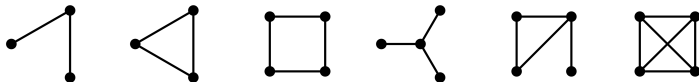
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# Goodness-of-fit using network motifs



- Network motifs have sociological interpretation (e.g. triangles).
- The first moments  $\mathbb{E}N(m)$ ,  $\mathbb{V}N(m)$  of the count are known under SBM [Picard *et al.* (2008)]:









$$\mathbb{E}_{SBM}N(m) \propto \mu_{SBM}(m) = f(\theta_{SBM})$$

- Motif probability can be estimated as

$$\hat{\mu}(m) = \sum_k Q_K^*(K) \tilde{\mathbb{E}}(\mu_{SBM}(m) | X, K)$$

→ Goodness of fit criterion

# Network motifs in the blogosphere

Motif	Count ( $\times 10^3$ )	Mean ( $\times 10^3$ )	Std. dev. ( $\times 10^3$ )	approx $p$ -value
	29.7	39.7	8.3	0.89
	3.8	4.6	1.3	0.69
	608.7	968.3	336.8	0.86
	279.8	428.9	154.0	0.83
	47.4	74.5	35.1	0.77
	270.5	397.0	177.0	0.75
	62.1	87.8	47.4	0.67
	6.5	8.8	5.4	0.61

No specific structure seems to be exceptional wrt the model's expectations.

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- SBM can be easily generalized to handle weighted graphs and covariates.
- SBM can be used as a proxy for smoother models, such as  $W$ -graphs.

- AIROLDI, E. M., COSTA, T. B. and CHAN, S. H. (2013). Stochastic blockmodel approximation of a graphon: Theory and consistent estimation. In *Advances in Neural Information Processing Systems*, 692–700.
- BEAL, J., M. and GHAHRAMANI, Z. (2003). The variational Bayesian EM algorithm for incomplete data: with application to scoring graphical model structures. *Bayes. Statist.* **7** 543–52.
- BOLLOBÁS, B., JANSON, S. and RIORDAN, O. (2007). The phase transition in inhomogeneous random graphs. *Rand. Struct. Algo.* **31** (1) 3–122.
- CERESSE, A., DAUDIN, J.-J. and PIERRE, L. (2012). Consistency of maximum-likelihood and variational estimators in the stochastic block model. *Electron. J. Statis.* **6** 1847–99.
- CHATTERJEE, S. (2012). Matrix estimation by Universal Singular Value Thresholding. Technical report, arXiv:1212.1247.
- DAUDIN, J.-J., PICARD, F. and ROBIN, S. (Jun, 2008). A mixture model for random graphs. *Stat. Comput.* **18** (2) 173–83.
- DAUDIN, J.-J., PIERRE, L. and VACHER, C. (2010). Model for heterogeneous random networks using continuous latent variables and an application to a tree–fungus network. *Biometrics.* **66** (4) 1043–1051.
- DEMPSTER, A. P., LAIRD, N. M. and RUBIN, D. B. (1977). Maximum likelihood from incomplete data via the EM algorithm. *J. R. Statist. Soc. B.* **39** 1–38.
- DIACONIS, P. and JANSON, S. (2008). Graph limits and exchangeable random graphs. *Rend. Mat. Appl.* **7** (28) 33–61.
- GAIL, S., DAUDIN, J.-J. and ROBIN, S. (2012). Accuracy of variational estimates for random graph mixture models. *Journal of Statistical Computation and Simulation.* **82** (6) 849–862.
- GHIMAN, M. and NEWMAN, M. E. J. (2002). Community strucutre in social and biological networks. *Proc. Natl. Acad. Sci. USA.* **99** (12) 7821–6.
- GOUDA, A. and SZÁNTAI, T. (2010). On numerical calculation of probabilities according to Dirichlet distribution. *Ann. Oper. Res.* **177** 185–200. DOI: 10.1007/s10479-009-0601-9.
- HANCOCK, M., RAFTERY, A. and TANTRUM, J. (2007). Model-based clustering for social networks. *JRSSA.* **170** (2) 301–54. doi: 10.1111/j.1467-985X.2007.00471.x.
- HOFING, J. A., MADIGAN, D., RAFTERY, A. E. and VOLINSKY, C. T. (1999). Bayesian model averaging: A tutorial. *Statistical Science.* **14** (4) 382–417.

- HOF, P. D., RAFTERY, A. E. and HANDCOCK, M. S. (2002). Latent space approaches to social network analysis. *J. Amer. Statist. Assoc.* **97** (460) 1090–98.
- JANKOLA, T. (2000). *Advanced mean field methods: theory and practice*. chapter Tutorial on variational approximation methods. MIT Press.
- LAPOUCHE, P., BIRMELE, E. and AMBROISE, C. (2012). Variational bayesian inference and complexity control for stochastic block models. *Statis. Model.* **12** (1) 93–115.
- LAPOUCHE, P. and ROBIN, S. (2013). Bayesian model averaging of stochastic block models to estimate the graphon function and motif frequencies in a W-graph model. Technical report, arXiv:1310.6150.
- LOVASZ, L. and SZEGEDY, B. (2006). Limits of dense graph sequences. *Journal of Combinatorial Theory, Series B.* **96** (6) 933 – 957.
- VON LUXBURG, U., BELKIN, M. and BOUSQUET, O. (2008). Consistency of spectral clustering. *Ann. Stat.* **36** (2) 555–586.
- MARADASSOU, M., ROBIN, S. and VACHER, C. (2010). Uncovering structure in valued graphs: a variational approach. *Ann. Appl. Statist.* **4** (2) 715–42.
- MARADASSOU, M. and MATIAS, C. (2015). Convergence of the groups posterior distribution in latent or stochastic block models. *Bernoulli*. ??–?? to appear.
- MATIAS, CATHERINE and ROBIN, STÉPHANE. (2014). Modeling heterogeneity in random graphs through latent space models: a selective review. *ESAIM: Proc.* **47** 55–74.
- NEWMAN, M. and GIRVAN, M. (2004). Finding and evaluating community structure in networks,. *Phys. Rev. E.* **69** 026113.
- NEWMAN, M. E. J. (2004). Fast algorithm for detecting community structure in networks. *Phys. Rev. E* (69) 066133.
- NOWICKI, K. and SNIJERS, T. (2001). Estimation and prediction for stochastic block-structures. *J. Amer. Statist. Assoc.* **96** 1077–87.
- PARISI, G. (1988). *Statistical Field Theory*. Addison Wesley, New York),.
- PICARD, F., DAUDIN, J.-J., KOSKAS, M., SCHBATH, S. and ROBIN, S. (2008). Assessing the exceptionality of network motifs,. *J. Comp. Biol.* **15** (1) 1–20.
- PICARD, F., MIELE, V., DAUDIN, J.-J., COTTRET, L. and ROBIN, S. (2009). Deciphering the connectivity structure of biological networks using mixnet. *BMC Bioinformatics. Suppl 6* S17. doi:10.1186/1471-2105-10-S6-S17.



VACHER, C., PIOU, D. and DESPREZ-LOUSTAU, M.-L. (2008). Architecture of an antagonistic tree/fungus network: The asymmetric influence of past evolutionary history. *PLoS ONE*. **3** (3) 1740. e1740. doi:10.1371/journal.pone.0001740.



VOISANT, S., MAGNIETTE, M.-L. M. and ROBIN, S. (2012). Variational bayes approach for model aggregation in unsupervised classification with markovian dependency. *Comput. Statis. & Data Analysis*. **56** (8) 2375 – 2387.



VANSWRIGHT, M. J. and JORDAN, M. I. (2008). Graphical models, exponential families, and variational inference. *Found. Trends Mach. Learn.* **1** (1–2) 1–305. <http://dx.doi.org/10.1561/22000000001>.