Bayesian modelling of brain network data via latent space models

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- Non-invasive imaging technologies provide accurate data on brain activity and structure at increasing resolution for multiple subjects
- Neuro-imaging study comprising data for m = 21 individuals (Landman et al., 2011).
- n = 68 brain regions (nodes), spatially located



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 Goal: investigate network connectivity patterns, accounting for anatomical constraints and unobservable patters (e.g. shapes, functionalities)

Brain Network Data

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Brain Network Data

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- a_{ij}^(k) = a_{ji}^(k) = 1 if at least one white matter fiber has been observed between regions i = 2,..., n and j = 1,...,i-1

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$$a_{ij}^{(k)} = a_{ji}^{(k)} = 0$$
 otherwise.





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- ► $a_{ij}^{(k)} = a_{ji}^{(k)} = 0$ otherwise. Anatomical information
- ► Spatial coordinates for the *i*-th region (x_i, y_i, z_i)
- Lobes and hemisphere membership
 - \rightarrow lobe_{ij} = 1 region *i* and region *j* are in the same lobe
 - \rightarrow hemi_{ij} = 1 region *i* and region *j* are in the same hemisphere





Latent Space Models - intuition



Developed in social sciences (e.g. Hoff et al., 2002; Hoff, 2008)



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Benefits

- ► Reduce dimensionality from n × (n − 1)/2 to n × H
- Takes into account network properties (e.g. transitivity, homophily)



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- ▶ Regions with similar propensities are more likely to be connected



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- Include covariates
 - → Connectivity as a function of anatomical constraints (distance, lobes)
- Estimate *local* clusters of brain regions
 - → Some regions might be similar only wrt *subset* of latent features.





Specification



• Focus on modelling
$$\mathbf{A} = \sum_{k=1}^{m} A^{(k)}$$

$$(a_{ij} \mid \pi_{ij}) \sim \text{Binom}(m, \pi_{ij})$$

 $\text{logit}(\pi_{ij}) = \beta_0 + \beta_1 \text{hem}_{ij} + \beta_2 \text{lobe}_{ij} + \beta_3 d_{ij} - \bar{d}_{ij}$

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$$d_{ij} = \sqrt{(x_i - x_j)^2 + (y_i - y_j)^2 + (z_i - z_j)^2}$$

 $\rightarrow\,$ Euclidean distance between region i and j in the original space

▶ $(\beta_1, \beta_2, \beta_3) \in \mathbb{R}^3$ effect of lobe and hemisphere membership and distance between regions

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 (β₁, β₂, β₃) ∈ ℝ³ effect of lobe and hemisphere membership and distance between regions

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$$\bar{d}_{ij} = \sqrt{(\bar{x}_i - \bar{x}_j)^2 + (\bar{y}_i - \bar{y}_j)^2 + (\bar{z}_i - \bar{z}_j)^2}$$

 \rightarrow Euclidean distance between region i and j in the latent space

• $(\bar{x}_i, \bar{y}_i, \bar{z}_i) \in \mathbb{R}^3$ latent coordinates of region *i*

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- Estimate groups of brain regions similar to *subsets* of latent features

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- Sparse Dirichlet on ν_x, ν_x and ν_y to favour deletion of redundant components (Rousseau and Mengersen, 2011)
- Gaussian for (β₀, β₁, β₂, β₃), conditionally conjugate trough the Pòlya-Gamma data augmentation (Polson et al., 2013).
- Metropolis step for updating the latent coordinates (Euclidean distance)



Results





Results 2



Inference on the separate partitions







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Inference on the separate partitions



...and on the coefficients

	Mean	Median	Std. Dev.	Cred. Int. _{95%}
Intercept	7.27	7.27	0.18	(6.94, 7.60)
hemisphere	0.60	0.61	0.18	(0.29, 0.92)
lobes	0.24	0.24	0.06	(0.13,0.35)
distance	-0.35	-0.35	0.06	(-0.47,-0.23)





Computational Drawbacks



- ► Issue: Inference relying on Markov Chain Monte Carlo scales poorly
- CPU time
 - \rightarrow n = 68, 2 min ×1000 it.
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- Approximate Bayesian inference
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 - \rightarrow Variational Inference (e.g. Blei et al., 2017)
 - Variational inference is widely popular in the network-science literature (Gollini and Murphy, 2016; Salter-Townshend and Murphy, 2013)
 - ► The Euclidean distance requires several Taylor expansions of the complete data log-likelihood





A different specification



Latent Factor Model (Hoff, 2008)

 $(a_{ij} \mid \pi_{ij}) \sim \text{Binom}(m, \pi_{ij})$ $\text{logit}(\pi_{ij}) = \beta_0 + \beta_1 \text{hem}_{ij} + \beta_2 \text{lobe}_{ij} + \beta_3 d_{ij} + \tilde{d}_{ij}$

$$\bullet \quad \tilde{d}_{ij} = \psi_x \tilde{x}_i \tilde{x}_j + \psi_y \tilde{y}_i \tilde{y}_j + \psi_z \tilde{z}_i \tilde{z}_j$$

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Gaussian priors

$$\begin{split} \boldsymbol{\beta} &= (\beta_0, \beta_1, \beta_2, \beta_3)^{\mathsf{T}} \sim \mathsf{N}_4(0, \boldsymbol{\Sigma}_0), \qquad \boldsymbol{\Sigma}_0 = \mathsf{diag}(\sigma_0, \dots, \sigma_3), \\ \tilde{x}_i \sim \mathsf{N}(0, 1), \quad \tilde{y}_i \sim \mathsf{N}(0, 1), \quad \tilde{z}_i \sim \mathsf{N}(0, 1) \quad i = 1, \dots, n, \\ (\psi_x, \psi_y, \psi_x) \sim \mathsf{N}_3(0, \gamma_{\psi_0} \boldsymbol{I}_3) \end{split}$$

• Conditional conjugancy introducing $(\omega_{ij} \mid -) \sim \mathsf{PG}(m, \mathsf{logit}(\pi_{ij}))$ 12



► Variational Bayes: find best approximation of the true posterior p in a restricted class Q of distributions

$$q^{\star}(\beta, \mathbf{W}, \omega, \psi) = \operatorname*{arg\,min}_{q \in \mathcal{Q}} \operatorname{KL} \left\{ q(\beta, \mathbf{W}, \omega, \psi) \mid \mid p(\beta, \mathbf{W}, \omega, \psi \mid \mathbf{A}) \right\}.$$



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► **Mean Field**: product restriction $Q = \{q(\beta, \mathbf{W}, \omega, \psi) : q(\beta, \mathbf{W}, \omega, \psi) = q(\beta)q(\mathbf{W})q(\omega)q(\psi)\}$



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- Analytical form for the optimal factors (same EF form than Full-Conditionals)

$$\begin{array}{l} q^{\star}(\boldsymbol{\beta}) \propto \exp\left\{\mathbb{E}_{q(\mathsf{W},\boldsymbol{\omega},\boldsymbol{\psi})}\log p(\boldsymbol{\beta}\mid -)\right\} & (\text{Gaussian}) \\ q^{\star}_{i}(\mathbf{w}_{i}) \propto \exp\left\{\mathbb{E}_{q(\boldsymbol{\beta},\mathsf{W}_{-i},\boldsymbol{\omega},\boldsymbol{\psi})}\log p(\mathbf{w}_{i}\mid -)\right\}, \quad i=1,\ldots,n & (\text{Gaussian}) \\ q^{\star}(\boldsymbol{\psi}) \propto \exp\left\{\mathbb{E}_{q(\boldsymbol{\beta},\mathsf{W},\boldsymbol{\omega})}\log p(\boldsymbol{\psi}\mid -)\right\} & (\text{Gaussian}) \\ q^{\star}_{ij}(\boldsymbol{\omega}_{ij}) \propto \exp\left\{\mathbb{E}_{q(\boldsymbol{\beta},\mathsf{w}_{i},\mathsf{w}_{j},\boldsymbol{\psi})}\log p(\boldsymbol{\omega}_{ij}\mid -)\right\} & (\text{Polya-Gamma}) \end{array}$$

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CAVI: cycle over each variational factor until convergence

Results, part II







- There has been considerable interest in Bayesian modelling of brain network data
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- Results suggests a general tendency of brain regions to connect with others that are spatially closer and belonging to the same lobe and hemisphere
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Thank you! Questions, comments?