

# Identifying Functional Co-Activation Patterns in Neuroimaging Studies Via Poisson Graphical Models

by: W Xue, J Kang, F DuBois Bowman, T Wager & J Guo  
*Biometrics* 70(4), 812-822



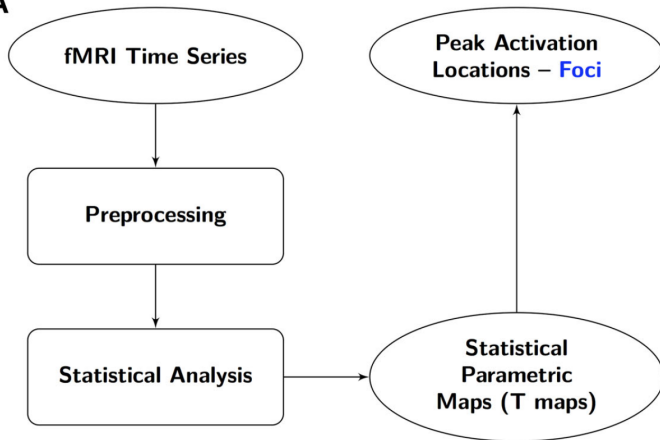
January 30, 2015

# Outline

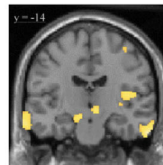
- 1 Motivation
- 2 Methods/Results
- 3 Conclusions

# fMRI experiments pipeline

A



x	y	z
24	-79	13
30	-83	23
18	-96	23
-16	-96	18
-22	-90	-17
-36	-93	14



# CBMA data & co-activation

**B**

Foci from Different Contrasts

Study	Contrast	Emotion	x	y	z
1	1	sad	-59	-14	-1
			...	...	...
			12	-74	-35
	2	happy	48	-78	-1
			...	...	...
			-57	-16	-2
2	3	anger	48	-74	-3
			...	...	...
			-24	0	-12
	4	sad	-57	-13	-2
			...	...	...
			14	-72	-32
	5	disgust	35	-80	15
			...	...	...
			-24	0	-12
...	....	...	...	...	...
164	437	fear	37	26	26
...	...	...	...	...	...



Regional Foci Count

Contrast	Region 1	Region 2	...	Region 19
1	0	1	...	1
2	1	2	...	1
3	1	0	...	1
4	3	2	...	5
5	2	0	...	3
...	....	....	...	....
437	2	1	...	2

# The Bivariate Poisson Model

- $i, j$  are 2 regions in brain,  $k$  denotes contrasts
- $X_{i,k}, X_{j,k}$  the number of foci
- When  $(X_{i,k}, X_{j,k}) \sim \text{BP}(\lambda_{ii}, \lambda_{jj}, \lambda_{ij})$  then:
  - $X_{i,k} \sim \text{Pois}(\lambda_{ii} + \lambda_{ij})$
  - $X_{j,k} \sim \text{Pois}(\lambda_{jj} + \lambda_{ij})$
  - Pmf is:

$$P(X_{i,k} = x_{i,k}, X_{j,k} = x_{j,k}) \\ = e^{-(\lambda_{ii} + \lambda_{jj} + \lambda_{ij})} \frac{\lambda_{ii}^{x_{i,k}} \lambda_{jj}^{x_{j,k}}}{x_{i,k}! x_{j,k}!} \sum_{s=0}^{\min(x_{i,k}, x_{j,k})} \binom{x_{i,k}}{s} \binom{x_{j,k}}{s} s! \left( \frac{\lambda_{ij}}{\lambda_{ii} \lambda_{jj}} \right)^s$$

# Network detection

- Parameter  $\boldsymbol{\lambda} = (\lambda_{ii}, \lambda_{jj}, \lambda_{ij})'$  fully describes network
- Covariance parameter  $\lambda_{ij}$  controls strength of co-activation
- A penalized likelihood framework is adopted:

$$\ell(\boldsymbol{\lambda}; \mathbf{X}_i, \mathbf{X}_j) - \theta \lambda_{ij}$$

- $\theta$  imposes sparsity on the network

# Latent variable representation

- The likelihood of the model is computationally demanding
- Let  $Y_{ij,k}$  be the total co-activations
- An alternative representation is:

$$X_{i,k} = Y_{ii,k} + Y_{ij,k}$$

$$X_{j,k} = Y_{jj,k} + Y_{ij,k}$$

- The complete model likelihood is then:

$$\ell_{\text{comp}}(\boldsymbol{\lambda}; \mathbf{Y}_{ij}, \mathbf{X}_i, \mathbf{X}_j) - \theta \lambda_{ij}$$

- Now, the EM of Karlis (2003) can be used for estimation

# Bivariate EM

- **E-step:**

$$\begin{aligned} Y_{ij,k}^{(t+1)} &= E[Y_{ij,k} | X_{i,k}, X_{j,k}; \boldsymbol{\lambda}^{(t)}] \\ &= \sum_{y_{ij,k}=0}^{\min(x_{i,k}, x_{j,k})} \frac{y_{ij,k} P(Y_{ij,k}, X_{i,k}, X_{j,k}; \boldsymbol{\lambda}^{(t)})}{\sum_{y_{ij,k}=0}^{\min(x_{i,k}, x_{j,k})} P(Y_{ij,k}, X_{i,k}, X_{j,k}; \boldsymbol{\lambda}^{(t)})} \end{aligned}$$

- **M-step:**

$$\begin{aligned} \lambda_{ij}^{(t+1)} &= \frac{\sum_{k=1}^n Y_{ij,k}^{(t+1)}}{\theta + n}, \\ \lambda_{ll}^{(t+1)} &= \frac{1}{n} \sum_{k=1}^n X_{l,k} - \frac{\theta + n}{n} \lambda_{ij}^{(t+1)} \quad \text{for } l = i, j. \end{aligned}$$



# The multivariate case

- Only 2-way interactions considered here
- Similar arguments as with 2D model:

$$X_{i,k} = \sum_{j=1}^p Y_{ij,k} \quad i = 1 \dots, p$$

- EM now minimizes:

$$\begin{aligned} & -l_{\text{comp}}(\lambda; \tilde{\mathbf{Y}}_1, \dots, \tilde{\mathbf{Y}}_n, \mathbf{X}_1, \dots, \mathbf{X}_n) + \theta \sum_{i=1}^p \sum_{j=i+1}^p \lambda_{ij} \\ & = \sum_{k=1}^n \sum_{i=1}^p \sum_{j=i}^p [\lambda_{ij} - Y_{ij,k} \log(\lambda_{ij})] + \theta \sum_{i=1}^p \sum_{j=i+1}^p \lambda_{ij}. \end{aligned}$$

# Choice of tuning parameter

- Optimal value of  $\theta$  is not known in practice
- Idea: split data in  $\mathbf{X}_{\text{train}}$ ,  $\mathbf{X}_{\text{test}}$  and use predictive log-likelihood:

$$l_{\text{obs}}(\hat{\boldsymbol{\lambda}}(\theta); \mathbf{X}_{\text{test}}) = \sum_{k=1}^n l_{\text{obs}}(\hat{\boldsymbol{\lambda}}(\theta); \mathbf{X}_{\text{test},k}).$$

- $n$ -fold cross validation for many  $\mathbf{X}_{\text{test}}$

# Testing significance

- 2 tests for significance of findings (non-zero covariances)
- Model detects networks but tests build on  $\lambda_{ij}^{\text{MLE}}$  (**no penalization**)
- Test I for pairs
  - $\mathcal{H}_0 : \lambda_{ij} = 0$  vs  $\mathcal{H}_1 : \lambda_{ij} > 0$
  - Contrast labels permuted for  $p$ -values
  - FDR applied
- Test II for full functional networks  $\Phi$ 
  - $\mathcal{H}_0 : \lambda_{ij} = 0, \forall \{i, j\} \in \Phi$  vs  $\mathcal{H}_1 : \exists \{i, j\} \in \Phi : \lambda_{ij} > 0$
  - Same permutation procedure as before

## Simulation studies 1/3

- Setup 1: 3 regions, 300 datasets, 100 bootstrap replicates for s.e.:

$$\lambda = \begin{bmatrix} 1 & 3 & 1 \\ & 2 & 5 \\ & & 3 \end{bmatrix}$$

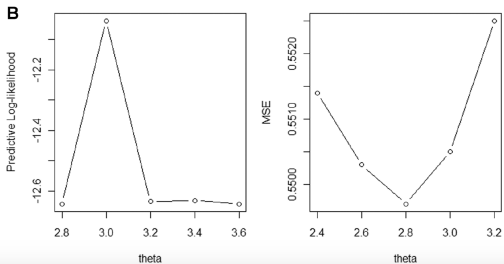
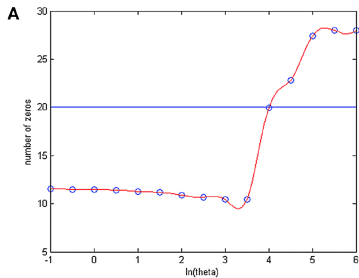
- Setup 2: 8 regions, 500 datasets:

$$\begin{array}{cccc} \lambda_{12} = 3 & \lambda_{15} = 4 & \lambda_{16} = 2 & \lambda_{27} = 2 \\ \lambda_{36} = 3 & \lambda_{48} = 4 & \lambda_{57} = 5 & \lambda_{78} = 1 \end{array}$$

# Simulation studies 2/3

Penalized multivariate Poisson model					
Bias (%)			Coverage rate		
0.0020 (0.20%)	0.0019 (0.06%)	0.0115 (1.15%)	93.33%	95.00%	90.67%
	0.0142 (0.71%)	0.0024 (0.05%)		96.00%	95.00%
		0.0169 (0.56%)			95.00%
Covariance method					
Bias (%)			Coverage rate		
0.1657 (16.57%)	0.0624 (2.08%)	0.1381 (13.81%)	92.00%	92.33%	91.67%
	0.0576 (2.88%)	0.1457 (2.91%)		96.33%	91.67%
		0.2747 (9.16%)			92.00%

# Simulation studies 3/3

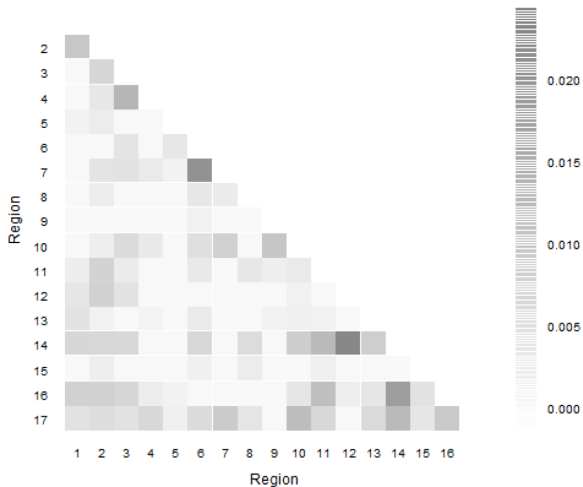


# Data

- 437 contrasts from 162 studies of emotion (Kober et al, 2008)
- On average, 6 foci per contrast
- GSK CIC atlas based on Harvard-Oxford atlas
- 19 ROIs in total
  - Dorsolateral prefrontal cortex reported w.p. 0.5 (highest)
  - Right globus pallidus reported w.p. 0.007 (lowest)
  - Others reported w.p. 0.140 (average)

## Results 1/4

- Emotional processing network detected: 17 ROIs, 79 connections

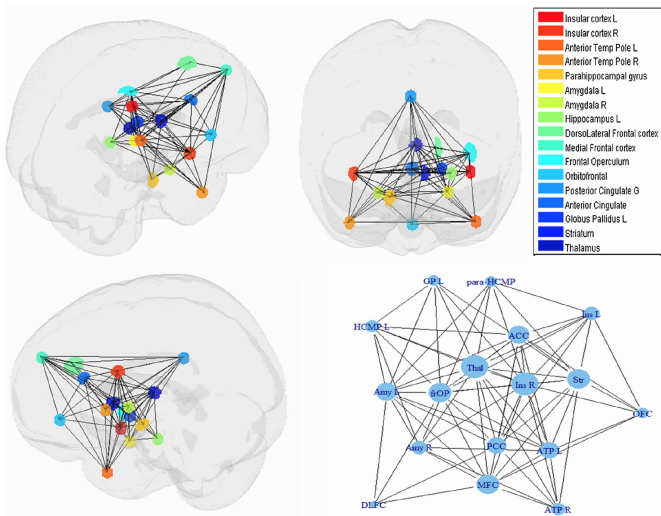




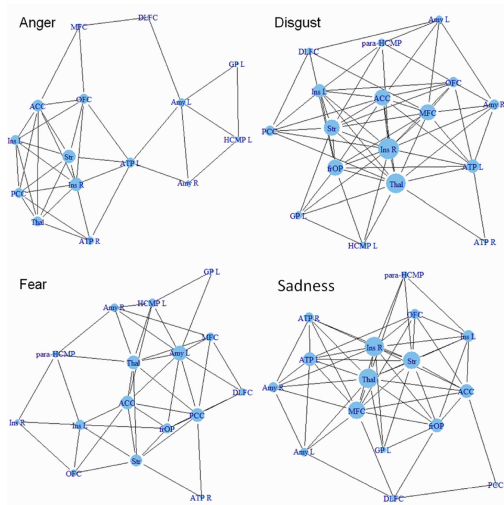
## Results 2/4

- Anterior cingulate cortex (ACC) most connections: 11
  - Orbitofrontal cortex  $\hat{\lambda}_{ij} = 0.023, p < 0.005$
  - Striatum  $\hat{\lambda}_{ij} = 0.018, p < 0.005$
  - Thalamus  $\hat{\lambda}_{ij} = 0.013, p < 0.005$
- Overall network significant as well  $p < 0.005$
- Clustering coef.  $C = 0.710$ , path length  $L = 1.129$
- Several regions with high degrees:
  - Right insular  $D = 14$
  - Thalamus  $D = 14$
  - Left amygdala  $D = 11$

# Results 3/4



# Results 4/4



# Discussion

- Pros:
  - New tool for CBMA data
  - Likelihood based
  - Nice intepretability
- Cons:
  - Brain tessellation may be subjective
  - No 3-way (or more) interactions
  - No voxel-wise rates

THANK YOU!!!

