

# Bayesian Nonparametric Models for Sparse Bipartite Networks

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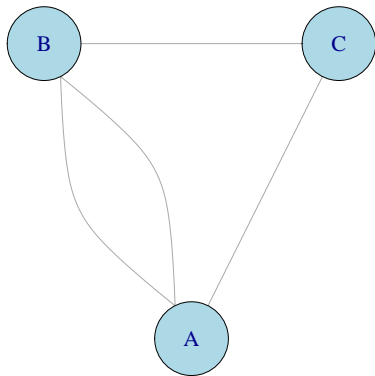
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Inria Center of University of Grenoble Alpes

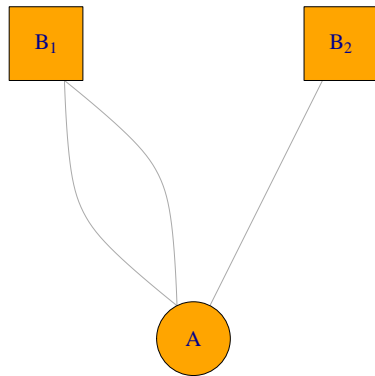
# Overview

- 1 Network Models and Exchangeability
- 2 Bayesian Nonparametrics (BNP)
- 3 Sparsity Results
- 4 Bayesian Model Estimation

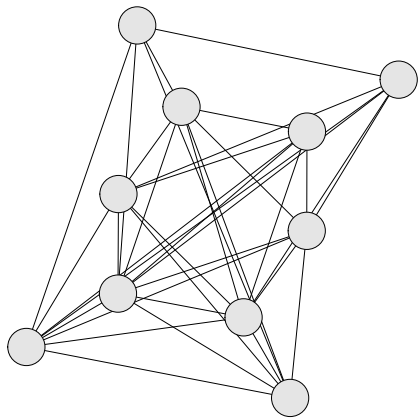
# Networks Structures



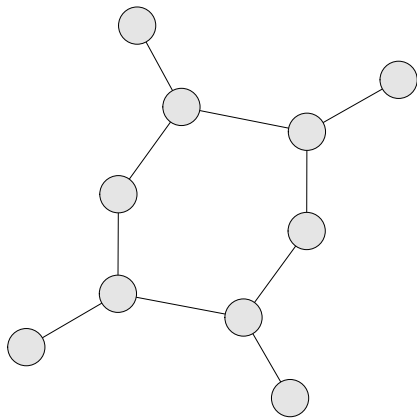
**Unipartite**



**Bipartite**



**Dense**



**Sparse**

# Definitions

For a **growing graph sequence**  $(\mathcal{Y}_n)_{n \in \mathbb{N}}$  (nodes and edges keep growing):

## Dense

The number of edges grows **at least quadratically** with the nodes.

$$e(\mathcal{Y}_n) = \Omega(v(\mathcal{Y}_n)^2)$$

## Sparse

The number of edges grows **sub-quadratically** with the nodes.

$$e(\mathcal{Y}_n) = o(v(\mathcal{Y}_n)^2)$$

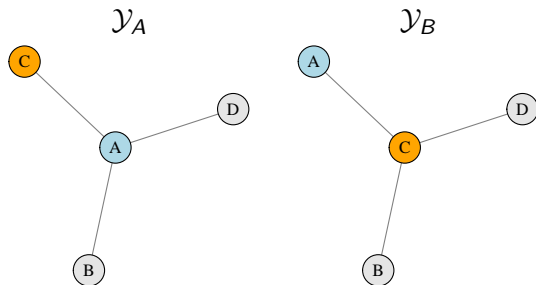
Equivalently, the sequence is **sparse** if the density converges to zero:

$$d(\mathcal{Y}_n) = \frac{e(\mathcal{Y}_n)}{v(\mathcal{Y}_n)^2} \xrightarrow{n \rightarrow \infty} 0.$$

Network models represent **probability distributions** over the space of possible graphs.

$$\Pr(\mathcal{Y}) = \Pr \left( \begin{array}{c} \text{C} \\ \diagdown \quad \diagup \\ \text{A} \\ \diagup \quad \diagdown \\ \text{B} \quad \text{D} \end{array} \right) = ?$$

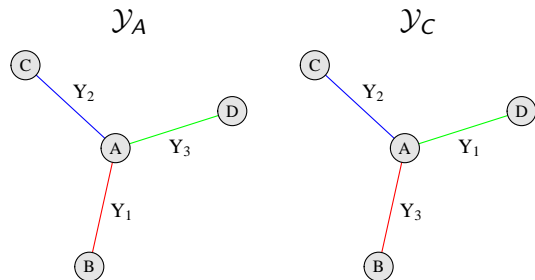
# Node Exchangeable Models



Most network models in the literature assume **node exchangeability**, i.e. **invariance** of the distribution under **permutation of nodes**. Reasonable when **node labels** are uninformative.

$$\Pr(\mathcal{Y}_A) = \Pr(\mathcal{Y}_B)$$

# Edge Exchangeable Models

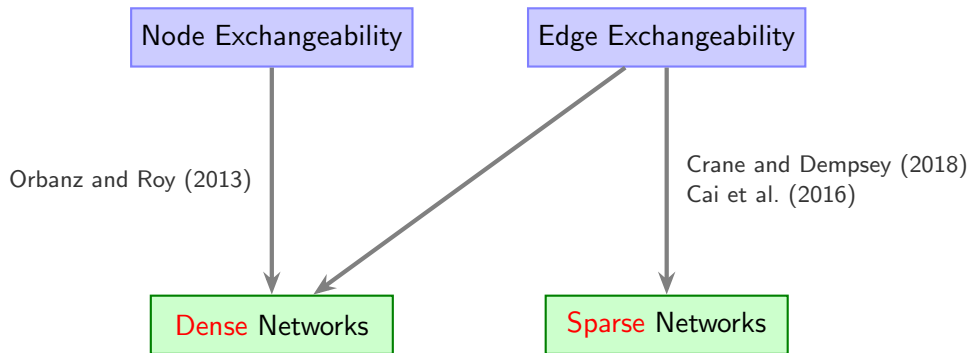


**Edge exchangeability** assumes **invariance** of the distribution under **permutation of edges** (the order of arrival).

$$\Pr(\mathcal{Y}_A) = \Pr(\mathcal{Y}_C)$$

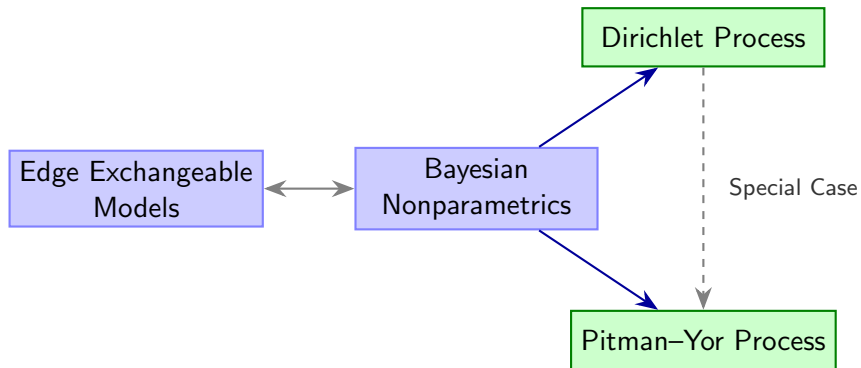


# Implications of Exchangeability



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# Dirichlet Process

The Dirichlet Process is a **distribution over probability distributions** (Ferguson, 1973).

$$G \sim \text{DP}(\alpha, G_0), \quad \alpha > 0$$
$$X_1, \dots, X_n \mid G \stackrel{\text{i.i.d.}}{\sim} G,$$

where  $\alpha$  is the **concentration** parameter, and  $G_0$  is a **probability distribution** (on the same space  $\Theta$  of the random variables  $X_i$ ).

**Property:**  $G$  is **discrete**  $\Rightarrow$  Samples from  $G$  contain **ties** with positive probability.

# Chinese Restaurant Process

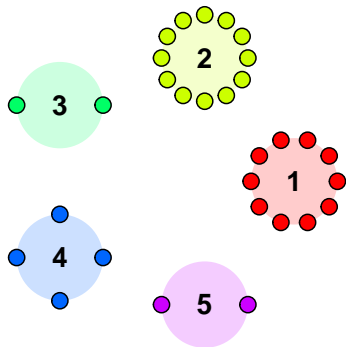
The probability of observing a new or an old value from a Dirichlet Process follows what is called the *Chinese Restaurant Process* (CRP).

## Dirichlet Process CRP

Let  $n_k$  be the frequency of the  $k$ -th unique value in the sample  $X_{1:n} = X_1, \dots, X_n$ .

$$\Pr(X_{n+1} = \text{Old } k\text{-th value} \mid X_{1:n}) = \frac{n_k}{\alpha + n}$$

$$\Pr(X_{n+1} = \text{New value} \mid X_{1:n}) = \frac{\alpha}{\alpha + n}$$



The Pitman-Yor Process is an extension of the DP (Pitman and Yor, 1997).

$$G \sim \text{PY}(\alpha, \sigma, G_0), \quad \sigma \in [0, 1)$$
$$X_1, \dots, X_n \mid G \stackrel{\text{i.i.d.}}{\sim} G.$$

**Connection:**  $\sigma = 0 \Rightarrow$  PY and DP coincide.

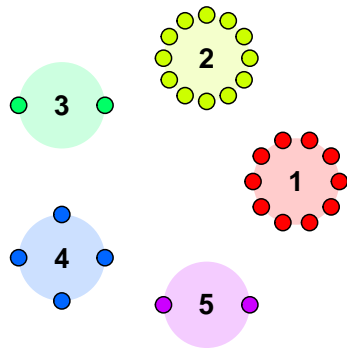
# Pitman-Yor CRP Construction

## Pitman-Yor Process CRP

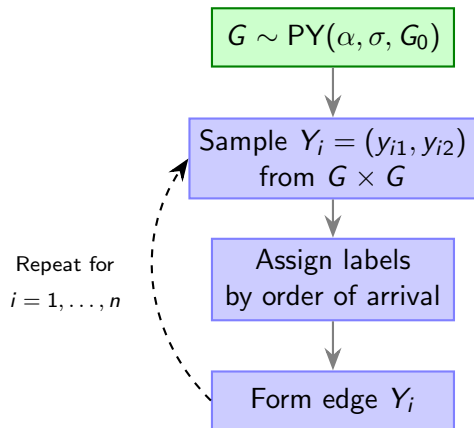
Let  $n_k$  be the frequency of the  $k$ -th unique value in the sample  $X_{1:n} = X_1, \dots, X_n$ , and let  $K_n$  be the number of unique values.

$$\Pr(X_{n+1} = \text{Old } k\text{-th value} \mid X_{1:n}) = \frac{n_k - \sigma}{\alpha + n}$$

$$\Pr(X_{n+1} = \text{New value} \mid X_{1:n}) = \frac{\alpha + \sigma K_n}{\alpha + n}$$

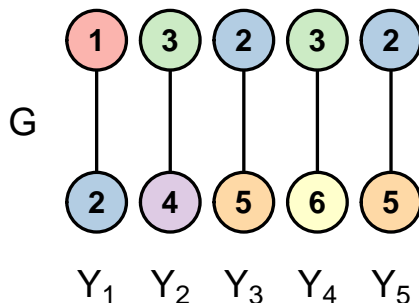


# BNP Generative Model: Unipartite Case



## Hierarchical Model

$$G \sim \text{PY}(\alpha, \sigma, G_0),$$
$$Y_1, \dots, Y_n \mid G \stackrel{i.i.d.}{\sim} G \times G$$





# BNP Generative Model: Bipartite Case

$$G_A \sim \text{PY}(\alpha_A, \sigma_A, G_0)$$

$$G_B \sim \text{PY}(\alpha_B, \sigma_B, G_0)$$

Sample  $Y_i = (y_{ia}, y_{ib})$   
from  $G_A \times G_B$

Assign labels  
by order of arrival

Form edge  $Y_i$

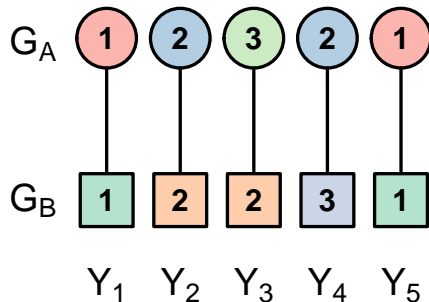
Repeat for  
 $i = 1, \dots, n$

## Hierarchical Model

$$G_A \sim \text{PY}(\alpha_A, \sigma_A, G_0)$$

$$G_B \sim \text{PY}(\alpha_B, \sigma_B, G_0)$$

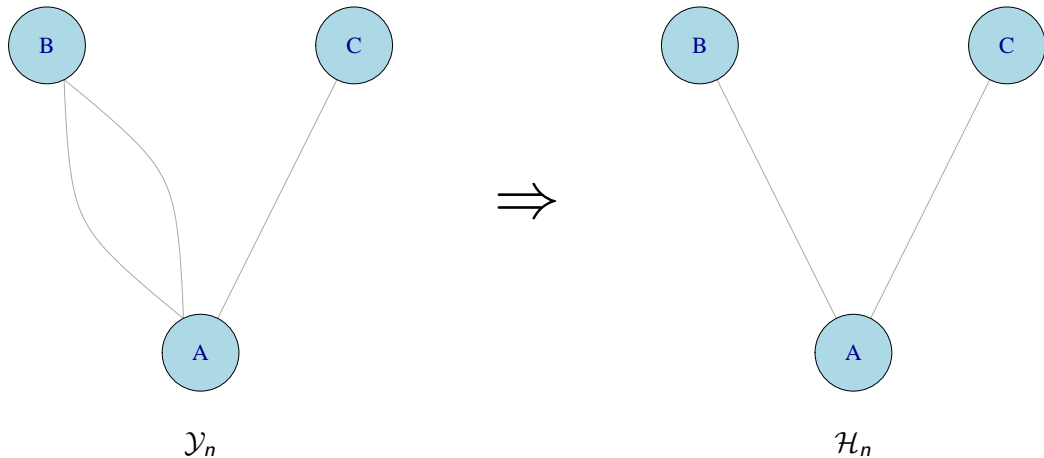
$$Y_1, \dots, Y_n \mid G_A, G_B \stackrel{i.i.d.}{\sim} G_A \times G_B$$



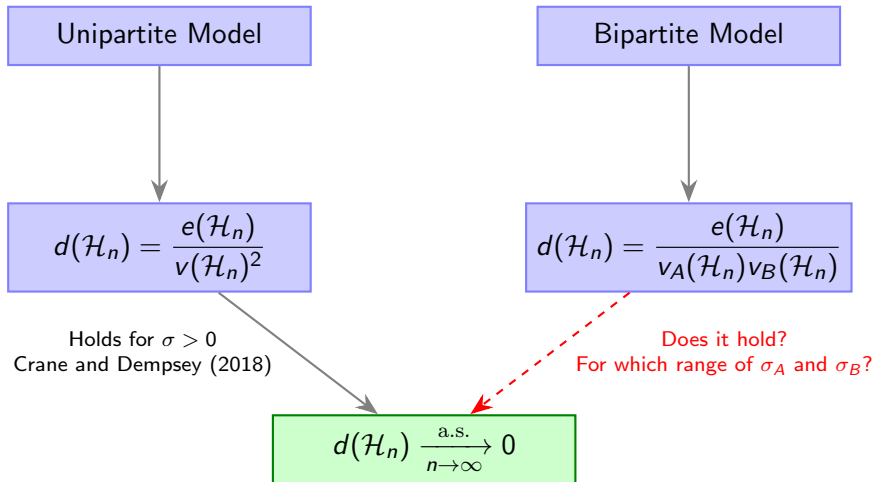
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# Binarisation



# Sparsity in Binary Graphs with PY Model



# Proposition: Bipartite PY Model is Sparse

## Proposition

The growing bipartite graph sequence  $(\mathcal{H}_n)_{n \in \mathbb{N}}$  produced by the PY process is sparse:

$$d(\mathcal{H}_n) = \frac{e(\mathcal{H}_n)}{v_A(\mathcal{H}_n)v_B(\mathcal{H}_n)} \xrightarrow[n \rightarrow \infty]{\text{a.s.}} 0,$$

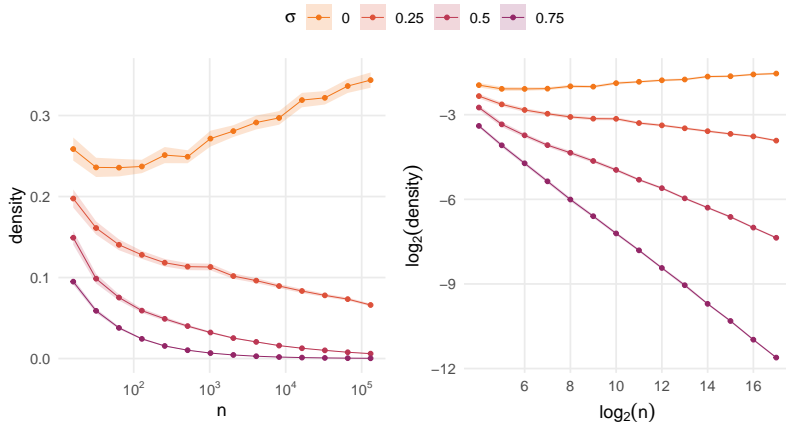
for either  $\sigma_A$  or  $\sigma_B$  in  $(0, 1)$ .

The proof technique follows closely that of Crane and Dempsey (2018) for the unipartite case.

### Remarks:

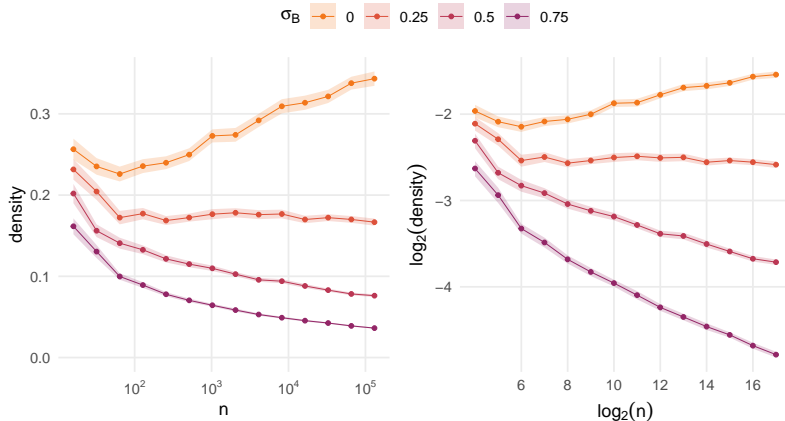
- If either  $\sigma_A = 0$  or  $\sigma_B = 0$ , sparsity is preserved (Asymmetric DP  $\times$  PY).
- If  $\sigma_A = \sigma_B = 0$ , sparsity is not guaranteed (Symmetric DP  $\times$  DP)

# Simulation Studies: Symmetric PY Model



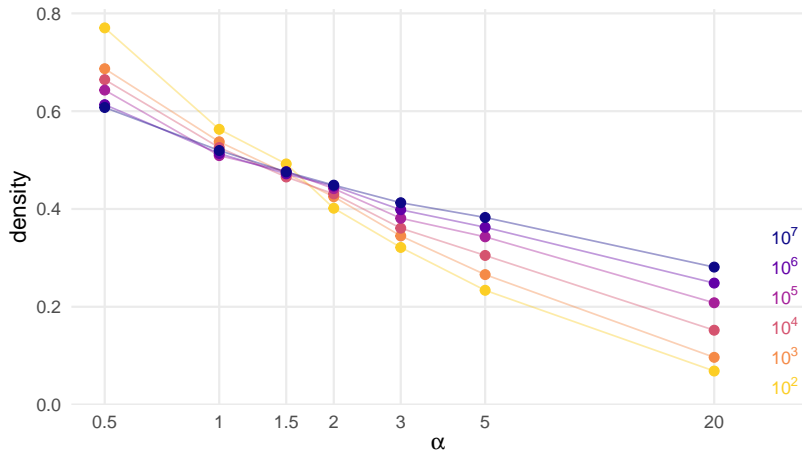
$$\text{PY}(5, \sigma_A) \times \text{PY}(5, \sigma_B), \sigma_A = \sigma_B = \sigma$$

# Simulation Studies: Asymmetric PY Model



$$\text{DP}(5) \times \text{PY}(5, \sigma_B), \sigma_A = 0$$

# Simulation Studies: Symmetric DP Model



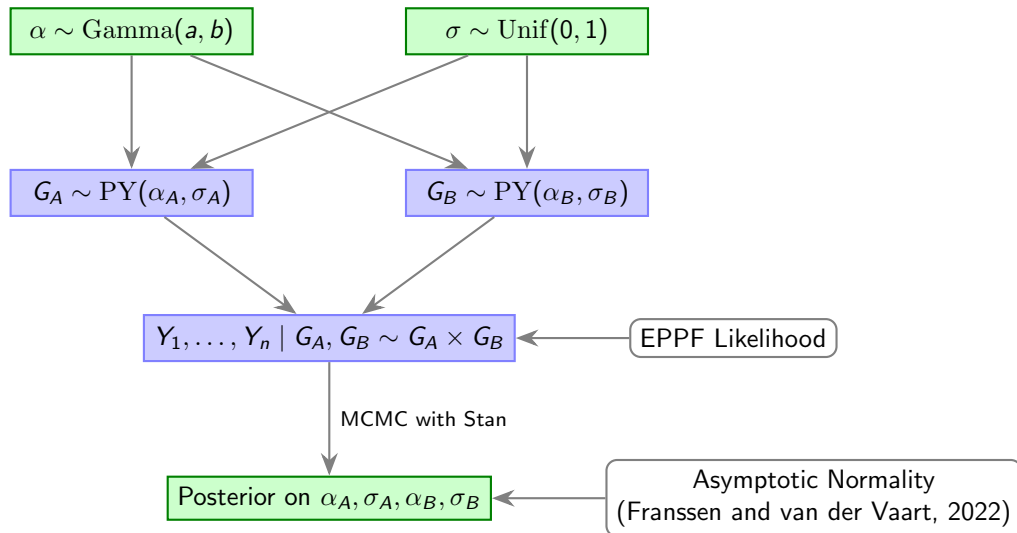
$$\text{DP}(\alpha_A) \times \text{DP}(\alpha_B), \alpha_A = \alpha_B = \alpha$$



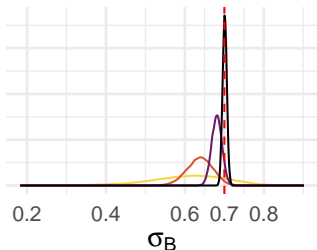
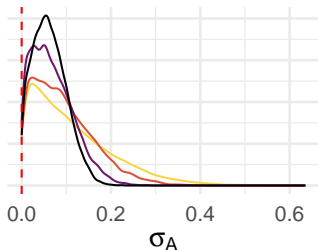
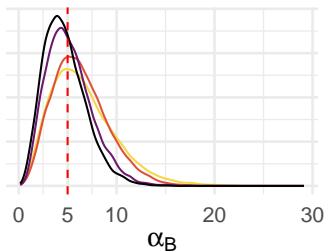
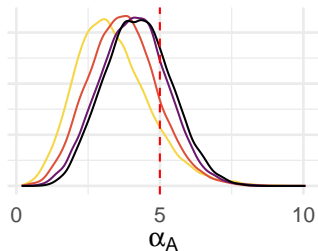
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# Bayesian Model



# Posterior Convergence: DP(5) $\times$ PY(5, 0.7)

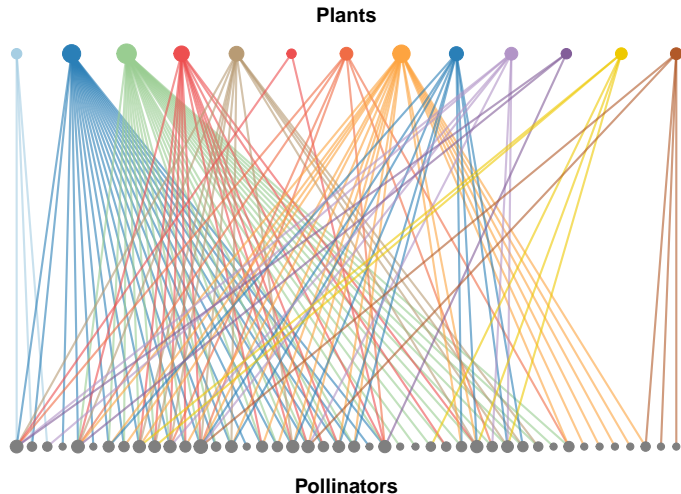


Edges  $10^2$   $10^3$   $10^4$   $10^5$

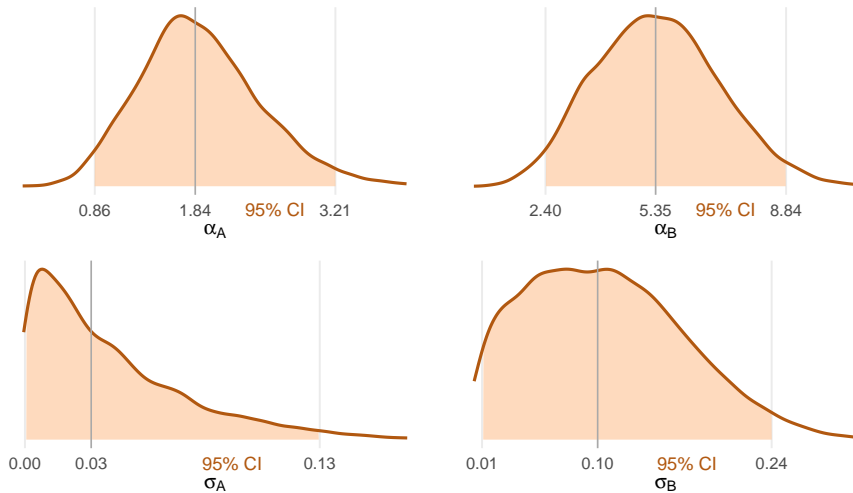
- Posterior distributions converge towards true parameter values (red dashed lines)
- Group B has a strong convergence for  $\sigma_B$
- Convergence is weaker in group A with  $\sigma_A = 0$

# Real Data

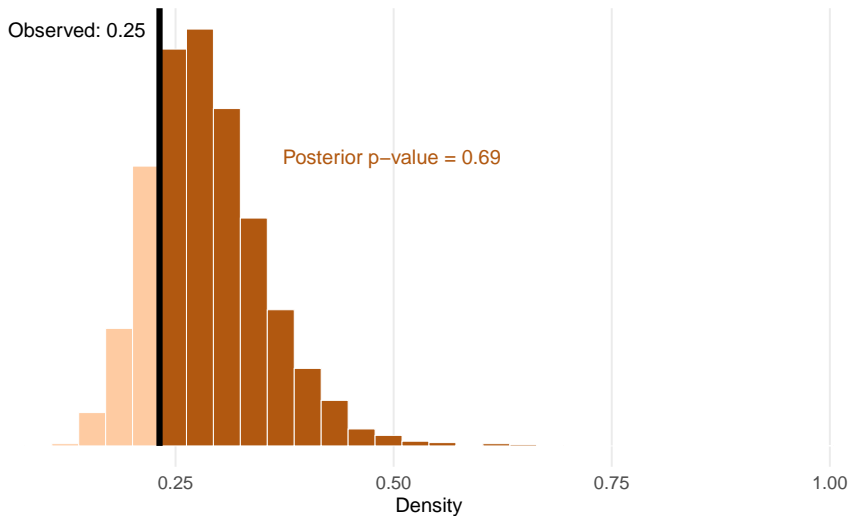
- Plants and Pollinators network data from forests in Piedmont North Carolina.
- Source: Web of Life. Original article: Motten (1986).
- 143 unique edges, 2225 observed interactions (sample size).
- 13 Plants (Group A)  
44 Pollinators (Group B).
- Density of 0.25.



# Real Data: Posteriors



# Real Data: Posterior Predictive Check



# Test for Sparsity

We want to formally test if one group is inducing sparsity.

$$H_0 : \sigma < \delta \quad H_1 : \sigma > \delta, \quad \text{for } \delta \text{ close to } 0.$$

The **Bayes Factor** tells us what is the strength of the evidence in favour of  $H_0$ , as a ratio of the probabilities of observing the data under the null over the alternative hypothesis:

$$\underbrace{\frac{p(\sigma \mid Y_{1:n}, H_0)}{p(\sigma \mid Y_{1:n}, H_1)}}_{\text{Posterior Odds}} = \underbrace{\frac{p(Y_{1:n} \mid \sigma, H_0)}{p(Y_{1:n} \mid \sigma, H_1)}}_{\text{Bayes Factor}} \times \underbrace{\frac{p(\sigma \mid H_0)}{p(\sigma \mid H_1)}}_{\text{Prior Odds}}.$$

# Bayes Factor on Real Data

$H_0 :$ $\delta$	$\sigma_A < \delta$		$\sigma_B < \delta$	
	BF	$\log_{10} \text{BF}$	BF	$\log_{10} \text{BF}$
0.01	26.69	1.43	3.55	0.55
0.05	47.54	1.68	5.33	0.73
0.10	117.90	2.07	8.85	0.95

Table: Bayes Factors for the discount parameters relative to threshold  $\delta$ .



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