

<sup>1</sup>Bocconi University, Milano

# Trees of random probability measures and Bayesian nonparametric modelling.

**Ascolani, F.** <sup>1</sup>    Lijoi, A. <sup>1</sup>    Prünster, I. <sup>1</sup>

July, 2022



A crucial question in Statistics is how to combine data from different sources:

- ▶ Homogeneity **within** each group.
- ▶ Heterogeneity **across** groups.

⇒ **partial exchangeability.**



Patients coming from **different hospitals.**

A crucial question in Statistics is how to combine data from different sources:

- ▶ Homogeneity **within** each group.
- ▶ Heterogeneity **across** groups.

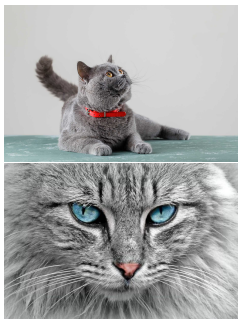
⇒ **partial exchangeability.**



Books belonging to the **same corpus.**

A crucial question in Statistics is how to combine data from different sources:

- ▶ Homogeneity **within** each group.
  - ▶ Heterogeneity **across** groups.
- ⇒ **partial exchangeability**.



Images of **similar subjects**.

# A first example

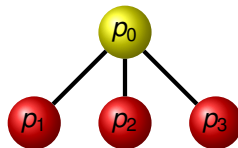


A popular model for incorporating heterogeneous information is given by **hierarchical Dirichlet processes**.

The model:

$$\begin{aligned} X_{i,j} | p_i &\stackrel{\text{i.i.d.}}{\sim} p_i, \\ p_i | p_0 &\sim \text{DP}(\theta, p_0), \\ p_0 &\sim \text{DP}(\theta, P_0), \end{aligned}$$

with  $P_0$  diffuse measure.



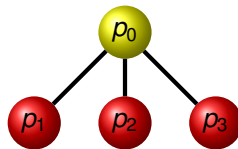
# A first example



A popular model for incorporating heterogeneous information is given by **hierarchical Dirichlet processes**.

The model:

$$\begin{aligned} X_{i,j} | p_i &\stackrel{\text{i.i.d.}}{\sim} p_i, \\ p_i | p_0 &\sim \text{DP}(\theta, p_0), \\ p_0 &\sim \text{DP}(\theta, P_0), \end{aligned}$$



with  $P_0$  diffuse measure.

- ▶ Each node is a **discrete** random measure.
- ▶ Often convolved with a suitable **kernel**  $k(y | x)$ .
- ▶ In **topic modelling**, the clusters correspond to different **topics**.

# A more complex problem...

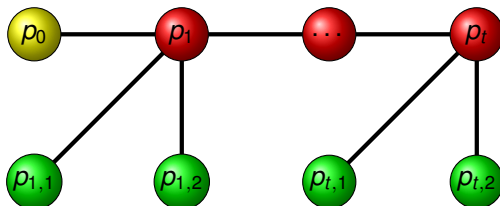


- ▶ *Data*: abstracts accepted at the NeurIPS conference from 2009 to 2019.
  - ▶ *Goal*: find suitable **topics** (i.e. probability distribution on words) that describe the abstracts.
  - ▶ *Problem*: incorporate the **temporal dynamics**.
- ⇒ hierarchical structure is no more appropriate!

...with a more complex structure.



- ▶  $p_0$  = **root** of the tree.
- ▶  $p_j$  = node associated to year  $2008 + j$ .
- ▶  $p_{j,i}$  = node associated to abstract  $i$  in year  $2008 + j$ .



⇒ **tree** structure!

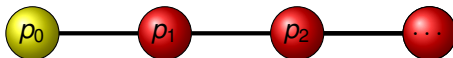




- ▶ A book can be seen as a **sequence of chapters**.
- ▶ Later chapters may have **similar topics** than the previous ones.
- ▶ How to introduce dependence between chapters?



- ▶ A book can be seen as a **sequence of chapters**.
- ▶ Later chapters may have **similar topics** than the previous ones.
- ▶ How to introduce dependence between chapters?



$\Rightarrow$  another special tree!



Many proposals for such structure:

- ▶ Extensions of Hierarchical Dirichlet processes ([Teh et al., 2006](#), [Caron et al., 2007, 2017](#))
- ▶ Other stick breaking priors ([Qi et al, 2008](#))
- ▶ Other classess of priors, e.g. Pólya trees ([Wang et al., 2021](#))



Many proposals for such structure:

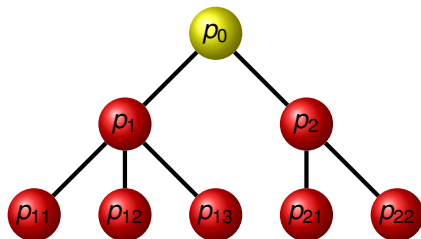
- ▶ Extensions of Hierarchical Dirichlet processes ([Teh et al., 2006](#), [Caron et al., 2007, 2017](#))
- ▶ Other stick breaking priors ([Qi et al, 2008](#))
- ▶ Other classess of priors, e.g. Pólya trees ([Wang et al., 2021](#))

What's new?

- ▶ **General framework** for constructing trees of random probability measures.
- ▶ Impact of the **shape of the tree**.
- ▶ Induced **clustering properties**.



The tree is described as follows:



LEVEL 0

LEVEL 1

LEVEL 2

- ▶  $X_{i,j}$  denotes the  $j$ -th observation at node  $i$  and  $X_{i,j} \mid p_i \stackrel{\text{i.i.d.}}{\sim} p_i$ .
- ▶ We call  $\text{MRCA}(i, j)$  the Most Recent Common Ancestor of nodes  $i$  and  $j$ .
- ▶ When talking about a specific level, we may omit the subscript  $k$ .



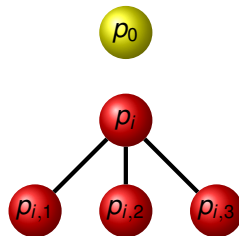
As a building block we consider Discrete RPMs, that is

$$p \stackrel{\text{a.s.}}{=} \sum_{k \geq 1} W_k \delta_{Z_k}, \quad \text{with} \quad \begin{cases} Z_k \stackrel{\text{i.i.d.}}{\sim} Q & \text{random atoms} \\ W_k & \text{random weights} \end{cases}$$

where  $Q$  is a probability distribution. We say  $p \sim \text{DRPM}(Q)$ .

1. **Root:**  $p_0 \sim \text{DRPM}(P_0)$ , with  $P_0$  diffuse measure.

2. **Edges:**  $p_{i,k} \mid p_i \stackrel{\text{i.i.d.}}{\sim} \text{DRPM}(p_i)$ .





It is a notion due to [Kingman \(1967\)](#).

## Definition

A random variable  $\mu$  is a **completely random measure** (CRM) if for any  $A_1, \dots, A_n$  measurable, with  $A_i \cap A_j = \emptyset$  for any  $i \neq j$ , the random variables  $\mu(A_1), \dots, \mu(A_n)$  are mutually independent.



It is a notion due to [Kingman \(1967\)](#).

## Definition

A random variable  $\mu$  is a **completely random measure** (CRM) if for any  $A_1, \dots, A_n$  measurable, with  $A_i \cap A_j = \emptyset$  for any  $i \neq j$ , the random variables  $\mu(A_1), \dots, \mu(A_n)$  are mutually independent.

Key property (**Lévy–Khintchine** representation):

$$E \left[ e^{-\lambda \mu(A)} \right] = e^{-\theta P_0(A) \psi(\lambda)}, \quad \psi(\lambda) = \int_{\mathbb{R}_+} (1 - e^{-\lambda s}) \rho(ds),$$

where  $\theta > 0$ ,  $P_0$  is a probability distribution and  $\rho$  is a measure on  $\mathbb{R}_+$  such that

$$\int_{\mathbb{R}_+} \min \{1, s\} \rho(ds) < \infty.$$





Under some technical conditions a CRM can be normalized (Regazzini et al. (2003)).

## Definition

Let  $\mu$  be a completely random measure, identified by  $(\rho, \theta, P_0)$ . Then  $p(\cdot) = \mu(\cdot)/\mu(\mathbb{X})$  is called **normalized random measure with independent increments** (NRMI) and we say  $p \sim \text{NRMI}(\rho, \theta, P_0)$ .



Under some technical conditions a CRM can be normalized  
(Regazzini et al. (2003)).

## Definition

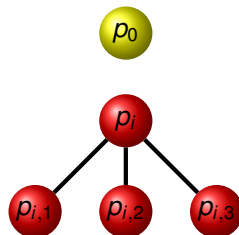
Let  $\mu$  be a completely random measure, identified by  $(\rho, \theta, P_0)$ .  
Then  $p(\cdot) = \mu(\cdot)/\mu(\mathbb{X})$  is called **normalized random measure with independent increments** (NRMI) and we say  $p \sim \text{NRMI}(\rho, \theta, P_0)$ .

Notable examples:

1. **Dirichlet process** (DP):  $\rho(ds) = s^{-1}e^{-s}ds$ .
2. **Normalized stable process** (NSP):  $\rho(ds) = \frac{\sigma}{\Gamma(1-\sigma)}s^{-1-\sigma}$ , with  $\sigma \in (0, 1)$ .

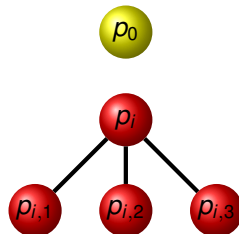
The model becomes:

1. **Root:**  $p_0 \sim \text{NRMI}(\rho, \theta, P_0)$ , with  $P_0$  diffuse measure.
2. **Edges:**  $p_{i,k} \mid p_i \stackrel{\text{i.i.d.}}{\sim} \text{NRMI}(\rho, \theta, p_i)$ .



The model becomes:

1. **Root:**  $p_0 \sim \text{NRMI}(\rho, \theta, P_0)$ , with  $P_0$  diffuse measure.
2. **Edges:**  $p_{i,k} \mid p_i \stackrel{\text{i.i.d.}}{\sim} \text{NRMI}(\rho, \theta, p_i)$ .



- ▶ Main advantage: **analytical tractability**, many prior and posterior results available.
- ▶ Probability of a tie:

$$\gamma = -\theta \int_{\mathbb{R}^+} u \left\{ \frac{d^2}{du^2} \psi(u) \right\} e^{-\theta \psi(u)} du \in (0, 1).$$



## Proposition

Let  $i$  and  $j$  be two nodes at level  $i$  and  $j$ , with MRCA at level  $k$ . Then

$$\text{Corr}(p_i(A), p_j(A)) = \frac{1 - (1 - \gamma)^{k+1}}{\sqrt{1 - (1 - \gamma)^{i+1}} \sqrt{1 - (1 - \gamma)^{j+1}}}$$

and

$$\text{Corr}(X_i, X_j) = 1 - (1 - \gamma)^{k+1}.$$

- ▶ As we move along the tree, the random measures become more and more correlated.
- ▶ As regards the DP and NSP we have:

$$\text{Corr}_{DP}(X_i, X_j) = 1 - \left( \frac{\theta}{1 + \theta} \right)^{k+1}, \quad \text{Corr}_{NSP}(X_i, X_j) = 1 - \sigma^{k+1}.$$



- Call  $K_n$  the number of distinct clusters in  $n$  observations. For simplicity, we focus on DP and NSP.

## Proposition

*Let  $k$  be a node at level  $k$  at which we collect  $n$  observations. Then*

$$K_{n,DP} \approx \underbrace{\log \dots \log n}_{k+1 \text{ times}}, \quad K_{n,NSP} \approx n^{\sigma^{k+1}},$$

*as  $n \rightarrow \infty$ .*



- Call  $K_n$  the number of distinct clusters in  $n$  observations. For simplicity, we focus on DP and NSP.

## Proposition

*Let  $k$  be a node at level  $k$  at which we collect  $n$  observations. Then*

$$K_{n,DP} \approx \underbrace{\log \dots \log n}_{k+1 \text{ times}}, \quad K_{n,NSP} \approx n^{\sigma^{k+1}},$$

*as  $n \rightarrow \infty$ .*

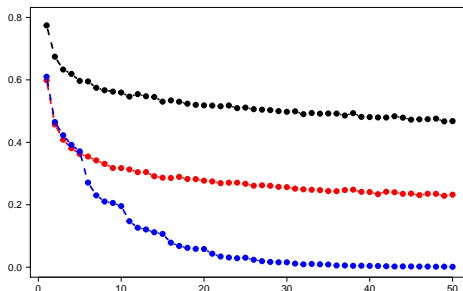
## Proposition

*Assume to collect  $m \geq 1$  observations at each level. Then*

$$\limsup K_{n,DP} < \infty, \quad \limsup K_{n,NSP} < \infty,$$

*as the number of levels diverges.*

- ▶ We can obtain the whole range of **clustering behaviours**.
- ▶ The **shape of the tree** is very relevant.



**Figure:** Average proportion of distinct values for groups of 20 observations.

In **black**: NSP with  $\sigma = 0.9$ .

In **red**: hierarchical NSP with  $\sigma = 0.9$ .

In **blue**: sequence of NSPs with  $\sigma = 0.9$ .





How are the observations generated at a generic level  $k$ ?

1. At level  $k$ ,  $n$  observations are sampled from

$$p_k \mid p_{k-1} \sim \text{NRMI}(\rho, \theta, p_{k-1}).$$

Since  $p_k \mid p_{k-1}$  is **discrete**, they are **grouped** in  $l_k \leq n$  clusters.



How are the observations generated at a generic level  $k$ ?

1. At level  $k$ ,  $n$  observations are sampled from  $p_k \mid p_{k-1} \sim \text{NRMI}(\rho, \theta, p_{k-1})$ .  
Since  $p_k \mid p_{k-1}$  is **discrete**, they are **grouped** in  $l_k \leq n$  clusters.
2. At level  $k - 1$ ,  $l_k$  observations are sampled from  $p_{k-1} \mid p_{k-2} \sim \text{NRMI}(\rho, \theta, p_{k-2})$ .  
Since  $p_{k-1} \mid p_{k-2}$  is **discrete**, they are **grouped** in  $l_{k-1} \leq l_k$  clusters.



How are the observations generated at a generic level  $k$ ?

1. At level  $k$ ,  $n$  observations are sampled from  $p_k \mid p_{k-1} \sim \text{NRMI}(\rho, \theta, p_{k-1})$ .  
Since  $p_k \mid p_{k-1}$  is **discrete**, they are **grouped** in  $l_k \leq n$  clusters.
2. At level  $k - 1$ ,  $l_k$  observations are sampled from  $p_{k-1} \mid p_{k-2} \sim \text{NRMI}(\rho, \theta, p_{k-2})$ .  
Since  $p_{k-1} \mid p_{k-2}$  is **discrete**, they are **grouped** in  $l_{k-1} \leq l_k$  clusters.
- $\vdots$
3. At level 0,  $l_1$  observations are sampled from  $p_0$ , with  $l_0 \leq l_1$  unique values from  $P_0$ .



How are the observations generated at a generic level  $k$ ?

1. At level  $k$ ,  $n$  observations are sampled from  $p_k \mid p_{k-1} \sim \text{NRMI}(\rho, \theta, p_{k-1})$ .  
Since  $p_k \mid p_{k-1}$  is **discrete**, they are **grouped** in  $l_k \leq n$  clusters.
  2. At level  $k - 1$ ,  $l_k$  observations are sampled from  $p_{k-1} \mid p_{k-2} \sim \text{NRMI}(\rho, \theta, p_{k-2})$ .  
Since  $p_{k-1} \mid p_{k-2}$  is **discrete**, they are **grouped** in  $l_{k-1} \leq l_k$  clusters.
  - $\vdots$
  3. At level 0,  $l_1$  observations are sampled from  $p_0$ , with  $l_0 \leq l_1$  unique values from  $P_0$ .
- ▶ All the unique values come from the root.
  - ▶ We have a **hidden clustering structure**. We only observe  $l_0$ .



In terms of the **Chinese Restaurant metaphor**:

- ▶ At each level the observations are subdivided in different **tables** (i.e. clusters).
- ▶ The **dishes** (i.e. unique values) come from the previous levels.
- ▶ Different tables may **share** the same dish.



In terms of the **Chinese Restaurant metaphor**:

- ▶ At each level the observations are subdivided in different **tables** (i.e. clusters).
- ▶ The **dishes** (i.e. unique values) come from the previous levels.
- ▶ Different tables may **share** the same dish.

Thus:

- ▶  $I_k$  = number of tables at level  $k$ .
- ▶ The levels share the same dishes.
- ▶ The root  $p_0$  becomes the **common menu**.



We consider a sample  $\mathbf{X} = \{X_{\mathbf{i},j}\}$ , with  $\mathbf{i}$  in the tree.

- ▶ Let  $X_1^*, \dots, X_r^*$  denote the distinct observations in sample  $\mathbf{X}$ .
- ▶ We call  $\mathbf{T}$  the **(latent) labels** of the tables.

For a fixed level  $k$ , we can then define

$l_{i,j}$  = number of tables at node  $i$  with dish  $j$ .

$q_{i,j,t}$  = number of customers at node  $i$  in table  $t$  eating dish  $j$ .

Conditional on  $\mathbf{T}$ , the posterior distribution becomes **accessible**!



Let  $\mathbf{U}$  be a positive random vector with density depending on  $\mathbf{T}$ .

## Theorem

We have

$$\mu_0 \mid (\mathbf{X}, \mathbf{T}, U_0) \stackrel{d}{=} \hat{\mu}_0 + \sum_{j=1}^r J_{0,j} \delta_{X_j^*},$$

where

1.  $\hat{\mu}_0$  is a CRM with intensity

$$\hat{\rho}_0(ds) = e^{-U_0 s} \rho(s) ds.$$

2. The  $J_{0,j}$ 's are independent and non-negative jumps with density

$$f_{0,j}(s \mid \mathbf{X}, \mathbf{T}) \propto s^{l_{1,j}} e^{-s U_0} \rho(s).$$





## Theorem

At level  $k$ , with ancestor  $p^*$ , we have

$$(\mu_{k,1}, \dots, \mu_{k,d}) \mid (p^*, \mathbf{X}, \mathbf{T}, \mathbf{U}) \stackrel{d}{=} (\hat{\mu}_1, \dots, \hat{\mu}_d) + \left( \sum_{j=1}^r \sum_{t=1}^{l_{1,j}} J_{1,j,t} \delta_{X_j^*}, \dots, \sum_{j=1}^r \sum_{t=1}^{l_{d,j}} J_{d,j,t} \delta_{X_j^*} \right),$$

where

1.  $\hat{\mu}_i$  is a CRM with baseline distribution  $p^*$  and intensity

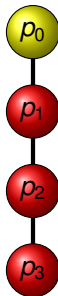
$$\hat{\rho}_i(ds) = e^{-U_i s} \rho(s) ds.$$

2. The  $J_{i,j,t}$ 's are independent and non-negative jumps with density

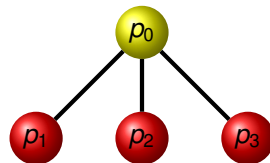
$$f_{i,j,t}(s \mid \mathbf{X}, \mathbf{T}) \propto s^{q_{i,j,t}} e^{-s U_i} \rho(s).$$

- ▶ We want to incorporate **chapters' specific information**.
- ▶ Two different structures.

Tree



Hierarchy



$p_i$  is the random measure associated to chapter  $i$ .



The **model**:

- ▶ Each  $p$  is a Dirichlet process, whose baseline distribution is given by the hierarchical structure.
- ▶ The hyperparameters at each node are endowed with vague priors.
- ▶ The root has the Dirichlet distribution as a baseline distribution, with common rate  $\alpha = 50/V$ .

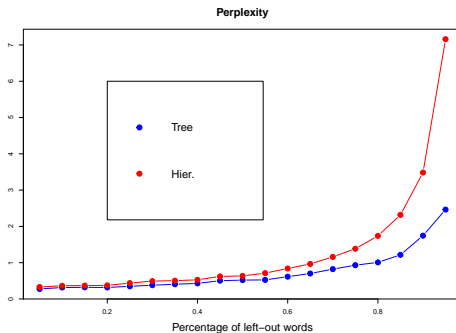


The **model**:

- ▶ Each  $p$  is a Dirichlet process, whose baseline distribution is given by the hierarchical structure.
- ▶ The hyperparameters at each node are endowed with vague priors.
- ▶ The root has the Dirichlet distribution as a baseline distribution, with common rate  $\alpha = 50/V$ .

The **data**:

- ▶ We consider the **initial three chapters**: standard pre-processing is applied.
- ▶ We randomly eliminate words from the second chapter and see whether the two methods are able to **recover** them.
- ▶ We measure goodness of fit in terms of the **perplexity** associated to the held-out words.



- ▶ The **lower** the perplexity the **better**.
- ▶ Results are averaged over 20 runs.
- ▶ **The tree always behaves better** and has good performances even with a high proportion of missing data.



## Summary:

- ▶ Many BNP models can be described using **trees**.
- ▶ If the nodes are given by NRMI, prior and posterior properties are available.
- ▶ We can use trees to make our learning process **explicit**.






## Summary:

- ▶ Many BNP models can be described using **trees**.
- ▶ If the nodes are given by NRMI, prior and posterior properties are available.
- ▶ We can use trees to make our learning process **explicit**.

## What's next?

- ▶ Construct a tree based on **covariates**.
- ▶ Study the asymptotic properties.
- ▶ Develop **efficient samplers** for posterior inference.



-  *Camerlenghi, F., Lijoi, A. and Prünster, I. (2019). Distribution theory for hierarchical processes. *Ann. Statist.* **47**, 67–92.*
-  *Kingman, J. F. C. (1967). Completely random measures. *Pacific J. Math.* **21**, 59–78.*
-  *Regazzini, E., Lijoi, A. and Prünster, I. (2003). Distributional results for means of normalized random measures with independent increments. *Ann. Statist.* **31**, 560–585.*