Discreteness and dependence: an effective interplay in Bayesian nonparametrics

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Outline

- ► DISCRETE PRIORS, RANDOM PARTITIONS AND PREDICTION
- ► PARTIAL EXCHANGEABILITY AND DEPENDENCE
- ► MULTIVARIATE SPECIES SAMPLING PROCESSES
- ► FLEXIBLE DEPENDENCE STRUCTURES & DISCRETENESS
 - ► CORRELATION STRUCTURE
 - ► RANDOM PARTITIONS
 - ► ATOMS
- ► MEASURING DEPENDENCE

Discrete priors, random

partitions and prediction

Exchangeable sequences

- ▶ Sequence $X = (X)_{n>1}$ of observations or latent features
- ► Standard assumption: exchangeability of X

$$\mathbf{X} \stackrel{\mathrm{d}}{=} \pi \mathbf{X} = (X_{\pi(i)})_{i \geq 1}$$
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 for any finite permutation π of $\mathbb N$

Conditional i.i.d. characterization

For any $n \geq 1$ and A_1, \ldots, A_n

$$\mathbb{P}[X_1 \in A_1, \dots, X_n \in A_n] = \int_{\mathsf{P}} \prod_{j=1}^n \rho(A_j) \ Q(\mathrm{d}\rho)$$

where P is the space of probability measures on the sample space or, equivalently,

$$X_i | \tilde{p} \stackrel{\text{iid}}{\sim} \tilde{p} \qquad \tilde{p} \sim Q$$

$$Q = \left\{ egin{array}{ll} & ext{probability measure on P} \\ & ext{de Finetti measure of } \textbf{\textit{X}} \ / \ ext{prior distribution} \end{array}
ight.$$

In B. de Finetti's own words

"Exchangeability" is the name and the notion proposed to replace "independent with unknown constant probability". Such terminology is clearly contradictory:
[...] let us think of drawing successively – with replacement – balls from an urn with unknown composition. Are the successive drawings independent?
They would be, for someone informed about the unknown (for other people) composition; for such a people the drawings are not informative, and the probability, for him, does not change. But, for any other observer, independence cannot, obviously, hold: the observed frequency is [...] informative ... [de Finetti (1979)]

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- Notion of "homogeneity" or "analogy" among observations for prediction
- ▶ In view of the exchangeability assumption, for any $m \ge 1$

$$\mathbb{P}[X_{n+1} \in A_1, \dots, X_{n+m} \in A_m \, | \, X_1, \dots, X_n] = \int_{P_{\mathbb{X}}} \prod_{i=1}^m p(A_i) \, \mathbf{Q}(\mathrm{d}p \, | \, X_1, \dots, X_n)$$

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► Most often Q selects discrete distributions

$$Q\Big(\Big\{p: \quad p=\sum_i \omega_i \delta_{\theta_i} \quad \text{for some } (\omega_i)_i \text{ and } (\theta_i)_i\Big\}\Big)=1$$

Species sampling

Species sampling process

- $\theta_i \stackrel{\text{iid}}{\sim} H \text{ with } H(\{x\}) = 0 \text{ for any } x$
- ▶ $(w_i)_{i \ge 1}$ non–negative and such that $\sum_i w_i \le 1$, almost surely
- $ightharpoonup (\theta_i)_{i\geq 1} \bot (w_i)_{i\geq 1}$

The random probability measure

$$\tilde{p} = \sum_{i \ge 1} w_i \, \delta_{\theta_i} + \left(1 - \sum_{i \ge 1} w_i\right) H$$

is a species sampling process (SSP). It is a proper SSP if $\sum_{i>1} w_i = 1$ (a.s.)

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Species sampling sequence

A sequence $(X_i)_{i>1}$ of random variables such that

$$X_i \mid \tilde{p} \stackrel{\text{iid}}{\sim} \tilde{p}$$
 & $\tilde{p} = \text{SSP}$

is termed a species sampling sequence (SSS).

A characterization

- $X_{1,n}^*,\ldots,X_{k_n,n}^*$ be the k_n distinct values in $X^{(n)}=(X_1,\ldots,X_n)$ with respective frequencies $\mathbf{n}=(n_1,\ldots,n_{k_n})$
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Theorem (Pitman, 1996)

A sequence $(X_n)_{n\geq 1}$ is a SSS if and only if there exist weights $\{p_{j,n}(n_1,\ldots,n_k):\ 1\leq j\leq k\leq n\}$ such that $X_1=\theta_1$ and for any $n\geq 1$

$$X_{n+1} \mid \boldsymbol{X}^{(n)} = \begin{cases} \theta_{n+1} & \text{with prob } p_{k_n+1,n}(\boldsymbol{n}^+) \\ X_{j,n}^* & \text{with prob } p_{k_n,n}(\boldsymbol{n}^{+j}) \end{cases}$$

where $\mathbf{n}^+=(n_1,\ldots,n_{k_n},1)$ and $\mathbf{n}^{+j}=(n_1,\ldots,n_j+1,\ldots,n_{k_n}).$

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- ▶ Predictive probability function $\implies \{p_{i,n}(n_1,\ldots,n_k): 1 \le j \le k+1, k \le n\}$
- ▶ If one is able to define a predictive probability function, then a SSS *X* is uniquely identified, without the need of specifying a prior *Q*. Very challenging a task!
- See Fortini, Ladelli & Regazzini (2002), Lee, Quintana, Müller & Trippa (2013).

Determining the prediction rule

- ▶ If $(X_n)_{n\geq 1}$ is a SSS, then $\mathbb{P}[X_i = X_j] > 0$ for any $i \neq j$.
- ▶ Random partition Ψ_n of $[n] = \{1, ..., n\}$, namely $i \sim j$ if and only if $X_i = X_j$

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Exchangeable partition probability function (EPPF)

$$\mathbb{P}[\Psi_n = \{C_1, \dots, C_k\}] = \Phi_k^{(n)}(n_1, \dots, n_k), \qquad n_i = \text{card}(C_i)$$

For example, with n = 4 and k = 2

$$\operatorname{Prob}(\underbrace{x_1}_{x_2}\underbrace{x_3}_{x_3}\underbrace{x_4}) = \Phi_2^{(4)}(\mathbf{n}_1 = \begin{bmatrix} 1 \\ 3 \end{bmatrix})$$

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Consistency condition & Symmetry

$$\Phi_k^{(n)}(n_1,\ldots,n_k) = \sum_{j=1}^k \Phi_k^{(n+1)}(n_1,\ldots,n_j+1,\ldots,n_k) + \Phi_{k+1}^{(n+1)}(n_1,\ldots,n_k,1)$$
 (*)

$$\Phi_k^{(n)}(n_1,\ldots,n_k) = \Phi_k^{(n)}(n_{\sigma(1)},\ldots,n_{\sigma(k)}) \quad \text{ for any finite permutation } \sigma \text{ of } [k] \quad (\Delta)$$

Random partitions and prediction

Prediction rule

The weights defined by

$$p_{j,n}(\mathbf{n}^{+j}) = \frac{\Phi_k^{(n+1)}(\mathbf{n}^{+j})}{\Phi_k^{(n)}(\mathbf{n})}, \qquad p_{k+1,n}(\mathbf{n}^+) = \frac{\Phi_{k+1}^{(n+1)}(\mathbf{n}^+)}{\Phi_k^{(n)}(\mathbf{n})}$$

identify the prediction rule of a SSS $(X_n)_{n>1}$.

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identify the prediction rule of a SSS $(X_n)_{n>1}$.

- ▶ If one can directly specify $\Phi_k^{(n)}$ that satisfies the conditions (*)–(Δ), then a SSS $(X_n)_{n\geq 1}$ is identified
- ► A prior Q is not involved in this construction, though it is uniquely identified by the EPPF $\{\Phi_k^{(n)}: 1 \le k \le n\}$
- ▶ Pro: assigning probability to observable events / quantities

An example: Gibbs-type priors (Gnedin & Pitman, 2006)

- ► Array of weights $\{V_{k,n}: 1 \le k \le n\}$ such that $V_{n,k} = V_{n+1,k+1} + (n-k\sigma)V_{n+1,k}$
- σ < 1
 </p>
- $\Phi_k^{(n)}(n_1,\ldots,n_k) = V_{n,k} \prod_{j=1}^k (1-\sigma)_{n_j-1}$, with $(a)_q = \Gamma(a+q)/\Gamma(a)$

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$$\begin{aligned} & p_{j,n}(\pmb{n}^{+j}) = \mathbb{P}[X_{n+1} = X_j^* \mid \pmb{X}^{(n)}] = \frac{V_{n+1,k}}{V_{n,k}}(n_j - \sigma) \\ & p_{k+1,n}(\pmb{n}^+) = \mathbb{P}[X_{n+1} = \text{"new"} \mid \pmb{X}^{(n)}] = \frac{V_{n+1,k+1}}{V_{n,k}} \end{aligned}$$

$$p_{k+1,n}(\mathbf{n}^+) = \mathbb{P}[X_{n+1} = \text{"new"} \mid \mathbf{X}^{(n)}] = \frac{v_{n+1,k+1}}{V_{n,k}}$$

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If K_n is the number of distinct values in X_1, \ldots, X_n and

$$c_n(\sigma) = \begin{cases} 1 & \sigma < 0 \\ \log n & \sigma = 0 \\ n^{\sigma} & \sigma \in (0, 1) \end{cases}$$

then

$$\frac{K_n}{c_n(\sigma)} \stackrel{\text{a.s.}}{\longrightarrow} S_{\sigma}$$

for random variable $S_{\sigma} > 0$

Dirichlet process: $\sigma = 0$ and $V_{n,k} = \theta^k/(\theta)_n$

$$\mathbb{P}[X_{n+1} = "new" \mid \mathbf{X}^{(n)}] = \frac{\theta}{\theta + n}$$

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with π a prior on the number of components. See De Blasi, L. and Prünster (2013) for some examples with different choices of π .

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- Gibbs-type priors and corresponding prediction rules introduced in BNP by L., Mena and Prünster (2007a, 2007b).
- ► Frequentist large sample properties in De Blasi, L. and Prünster (2013).
- ► Review in De Blasi et al. (2015)

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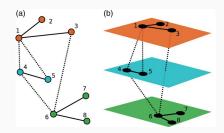
- Symmetry and, more importantly, consistency conditions are automatically satisfied in this case: no need to check their validity!
- Consistency is crucial for prediction and validity of inferential procedures. Lack of it, as in some finitely exchangeable models, only allows exploratory data analysis.

Partial exchangeability and

dependence

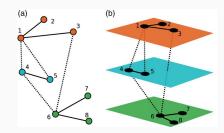
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- Not a realistic assumption in several applications



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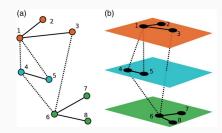
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- Edges between nodes in the same layer or across different layers
- Clustering of nodes based on connections

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- ► Edges between nodes in the same layer or across different layers
- Clustering of nodes based on connections
 - ► Bill co-sponsorship (*L* = political parties)
 - ► Criminal networks (*L* = main "clan" affiliation)

Examples:

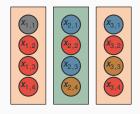
- ▶ ...
- Francesco Gaffi's talk yesterday & Filippo Ascolani's talk tomorrow

Beyond exchangeability

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Beyond exchangeability

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- Observations from different samples/groups
- reasonable dependence assumption
 - Homogeneity within each group
 - Heterogeneity across different groups
- lacktriangle Two levels of clustering: samples/groups $ilde{p}_j$ and observations $X_{i,j}$

Multiple samples and partial exchangeability

► Assumption: J samples $X_1 = (X_{1,i})_{i \ge 1}, \dots, X_J = (X_{J,i})_{i \ge 1}$ are partially exchangeable, i.e. for any finite permutations π_1, \dots, π_J of $\mathbb N$

$$(\boldsymbol{X}_1,\ldots,\boldsymbol{X}_J)\stackrel{\mathrm{d}}{=}(\pi_1\boldsymbol{X}_1,\ldots,\pi_J\boldsymbol{X}_J)$$

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With
$$J = 2$$
 samples (red & green):

$$(X_{1,1}, X_{1,2}, X_{2,1}) \stackrel{\mathrm{d}}{=} (X_{1,2}, X_{1,1}, X_{2,1})$$

$$(X_{1,1}, X_{1,2}, X_{2,1}) \stackrel{\mathrm{d}}{\neq} (X_{1,1}, X_{2,1}, X_{1,2})$$

Multiple samples and partial exchangeability

▶ **Assumption:** J samples $X_1 = (X_{1,i})_{i \ge 1}, \ldots, X_J = (X_{J,i})_{i \ge 1}$ are *partially exchangeable*, i.e. for any finite permutations π_1, \ldots, π_J of $\mathbb N$

$$(\boldsymbol{X}_1,\ldots,\boldsymbol{X}_J)\stackrel{\mathrm{d}}{=}(\pi_1\boldsymbol{X}_1,\ldots,\pi_J\boldsymbol{X}_J)$$

With J = 2 samples (red & green):

$$(X_{1,1}, X_{1,2}, X_{2,1}) \stackrel{d}{=} (X_{1,2}, X_{1,1}, X_{2,1})$$

 $(X_{1,1}, X_{1,2}, X_{2,1}) \stackrel{d}{\neq} (X_{1,1}, X_{2,1}, X_{1,2})$

Conditional independence characterization
$$(J=2)$$

$$\mathbb{P}\left[X_{1,1} \in A_1, \cdots, X_{1,N_1} \in A_{N_1}, \ X_{2,1} \in B_1, \cdots, X_{2,N_2} \in B_{N_2}\right]$$

$$= \int_{P_{\mathbb{X}}^2} \prod_{i=1}^{N_1} p_1(A_i) \prod_{j=1}^{N_2} p_2(B_j) \ Q_2(\mathrm{d}p_1, \mathrm{d}p_2)$$

$$(X_{1,i}, X_{2,j}) \mid (\tilde{p}_1, \tilde{p}_2) \stackrel{\mathrm{iid}}{\sim} \tilde{p}_1 \times \tilde{p}_2, \qquad (\tilde{p}_1, \tilde{p}_2) \sim Q_2$$

de Finetti (1938), Cifarelli & Regazzini (1978), MacEachern (1999, 2000), ...

Range of dependence

 $Q_2(dp_1, dp_2) = Q_1^*(dp_1) Q_2^*(dp_2)$

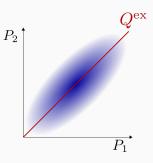
$$Q_2(\{p_1=p_2\})=1 \Longrightarrow$$

homogeneity across samples (maximal dependence)

unconditional independence (maximal heterogeneity)

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 \Longrightarrow homogeneity across samples (maximal dependence)
$$Q_2(\mathrm{d}p_1,\mathrm{d}p_2)=Q_1^*(\mathrm{d}p_1)\,Q_2^*(\mathrm{d}p_2) \Longrightarrow$$
 unconditional independence (maximal heterogeneity)



Intermediate cases: partial exchangeability

In this talk \tilde{p}_i 's are discrete

$$\tilde{p}_1 = \sum_{j \ge 1} w_{1,j} \, \delta_{Z_{1,j}} \qquad \tilde{p}_2 = \sum_{j \ge 1} w_{2,j} \, \delta_{Z_{2,j}}$$

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 $ightharpoonup k \le N_1 + N_2$ distinct values in the samples

$$extbf{X}_1^{(N_1)} = (X_{1,1}, \dots, X_{1,N_1}) \qquad extbf{X}_2^{(N_2)} = (X_{2,1}, \dots, X_{2,N_2})$$

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▶ $\mathbf{n}_1 = (n_{1,1}, \dots, n_{1,k})$ & $\mathbf{n}_2 = (n_{2,1}, \dots, n_{2,k})$ frequency vectors

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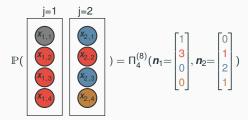
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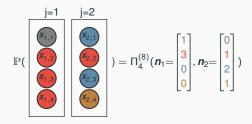
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- $ightharpoonup n_{1,j}n_{2,j} \neq 0$ whenever the jth distinct value is shared across the two samples
- ▶ Induced random partition Ψ_N of $[N] = \{1, ..., N_1 + N_2\}$, characterized by the partially exchangeable partition probability function

$$\mathsf{pEPPF} = \mathbb{P}[\Psi_n = \{C_1,\dots,C_k\}] = \Pi_k^{(N)}(\pmb{n}_1,\pmb{n}_2)$$
 where $\mathsf{card}(C_i) = n_{1:i} + n_{2:i} > 1.$

Joint distribution of the induced random partition:



Joint distribution of the induced random partition:



Marginal distribution of the single sample partition:

$$\mathbb{P}(\sqrt[k_{1,1}]{k_{1,2}}\sqrt[k_{1,3}]{k_{1,4}}) = \Phi_2^{(4)}(\textit{\textbf{n}}_1 = \begin{bmatrix} 1 \\ 3 \end{bmatrix})$$

Evaluating the pEPPF

▶ By definition

$$\Pi_k^{(N)}(\mathbf{n}_1,\mathbf{n}_2) = \mathbb{E} \sum_{\substack{i_1 \neq \cdots \neq i_k \\ j_1 \neq \cdots \neq j_k}} w_{1,i_1}^{n_{1,1}} \cdots w_{1,i_k}^{n_{1,k}} w_{2,j_1}^{n_{2,1}} \cdots w_{2,j_k}^{n_{2,j_k}}$$

which is impossible to evaluate

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which is impossible to evaluate

The most convenient form for actual evaluation is

$$\Pi_k^{(N)}(\textbf{\textit{n}}_1,\textbf{\textit{n}}_2) = \mathbb{E} \int_{\mathbb{X}_d^k} \prod_{j=1}^k \tilde{\rho}_1^{n_{1,j}}(\mathrm{d}x_j) \, \tilde{\rho}_2^{n_{2,k}}(\mathrm{d}x_k)$$

with closed form expressions available for models based on completely random measures

- ► GM-dependent NRMIs (L., Nipoti and Prünster, 2014)
- ► Hierarchical NRMIs (Camerlenghi et al., 2019)
- ► Latent nested processes (Camerlenghi et al., 2019)
- Bivariate Pitman–Yor process (Leisen and L., 2011)

Approaching dependence

In this talk: general framework

$$(\tilde{p}_1,\ldots,\tilde{p}_J)$$
 multivariate species sampling process

- ► Which dependence is induced among
 - ▶ the random probability measures \tilde{p}_i ?
 - the observations, within and between samples?
- Discreteness is helpful in shaping the dependence structure, which may be characterized through
 - correlation
 - the induced random partition
 - ► the atoms of the \tilde{p}_i 's

Multivariate species sampling

process

Multivariate species sampling processes

Definition

A vector of random probability measures $(\tilde{p}_1,\ldots,\tilde{p}_J)$ is a *multivariate species sampling process* (mSSP) if for any $j\in\{1,\ldots,J\}$

$$\tilde{p}_j = \sum_{h \ge 1} w_{j,h} \, \delta_{\theta_h} + \Big(1 - \sum_{h \ge 1} w_{j,h} \Big) H$$

- lacktriangledown θ_h are iid from the non–atomic probability measure H
- $\blacktriangleright (w_{1,h},...,w_{J,h})_{h\geq 1} \bot (\theta_h)_{h\geq 1}$

If $\sum_{h\geq 1} w_{j,h} = 1$, a.s., for every j, then $(\tilde{p}_1, \dots, \tilde{p}_J)$ is said *proper*.

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Multivariate species sampling sequence

The array of random elements $\{X_j = (X_{j,i})_{i \geq 1}: j = 1, ..., J\}$ is a *multivariate* species sampling sequence (mSSS) if

- ► it is partially exchangeable
- ▶ its de Finetti measure is a mSSP

A predictive characterization (with J = 2)

- Does there exist a predictive characterization that mimics the one seen for SSS?
- ► If yes, does the prediction rule relate to the induced random partition?

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Let $\theta_h \stackrel{\text{iid}}{\sim} H$ and X_1^*, \dots, X_k^* the k distinct values observed in the samples $\mathbf{X}_j^{(N_j)} = (X_{1,j}, \dots, X_{N_j,j})$, for j = 1, 2, with frequencies $\mathbf{n} = (\mathbf{n}_1, \mathbf{n}_2)$.

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- Does there exist a predictive characterization that mimics the one seen for SSS?
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Predictive characterization

The array of sequences of random variables (X_1, X_2) is a mSSS if an only if there exists weights $\{p_{i,j}: 1 \leq j \leq k+1, \ n \geq k\}$ such that $X_{1,1} = \theta_1$ and

$$X_{1,N+1} \mid (\boldsymbol{X}_1^{(N_1)}, \boldsymbol{X}_2^{(N_2)}) = \left\{ \begin{array}{ll} \theta_{N+1} & \quad \text{with prob } p_{1,k+1}(\boldsymbol{n}^+) \\ X_\ell^* & \quad \text{with prob } p_{1,\ell}(\boldsymbol{n}^{+\ell}) \end{array} \right.$$

where $N = N_1 + N_2$ and

$$\mathbf{n}^+ = (n_{1,1}, \dots, n_{k,1}, 1, \mathbf{n}_2), \quad \mathbf{n}^{+\ell} = (n_{1,1}, \dots, n_{\ell,1} + 1, \dots, n_{k,1}, \mathbf{n}_2)$$

Similar expression if the (N + 1)-th observation is from sample 2.

Prediction and random partition

- If one is able to determine a multivariate predictive probability function {p_{i,j}: 1 ≤ j ≤ k + 1 ≤ n + 1; i = 1,2} that satisfies some suitable conditions then a mSSS is identified (Franzolini et al., 2022+).
- ► Is there a connection between the prediction rule and the distribution of the partially exchangeable partition probability function (pEPPF) that extends what is known in the exchangeable case?
- ► Can one deduce a Pólya urn scheme from the pEPPF $\Pi_k^{(N)}$?

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$$\begin{split} \rho_{1,k+1}(\textbf{\textit{n}}) &= \frac{\Pi_{k+1}^{(N+1)}(\textbf{\textit{n}}_1^+,\textbf{\textit{n}}_2)}{\Pi_k^{(N)}(\textbf{\textit{n}}_1,\textbf{\textit{n}}_2)} = \mathbb{P}[X_{1,N_1+1} = \text{"new"} \, | \, \textbf{\textit{X}}_1^{(N_1)},\textbf{\textit{X}}_2^{(N_2)}] \\ \rho_{1,\ell}(\textbf{\textit{n}}) &= \frac{\Pi_k^{(N_1+1)}(\textbf{\textit{n}}_1^{+\ell},\textbf{\textit{n}}_2)}{\Pi_k^{(N)}(\textbf{\textit{n}}_1,\textbf{\textit{n}}_2)} = \mathbb{P}[X_{1,N_1+1} = X_\ell^* \, | \, \textbf{\textit{X}}_1^{(N_1)},\textbf{\textit{X}}_2^{(N_2)}] \end{split}$$

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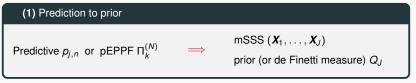
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► Marginally each X_i is a SSS, for i = 1, 2.

From prediction to prior & viceversa



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Approach (1)

- Assess probabilities related only to observable quantities/events
- ► Very challenging (not clear whether even possible) to implement!

From prediction to prior & viceversa

Predictive
$$p_{j,n}$$
 or pEPPF $\Pi_k^{(N)}$ \Longrightarrow $mSSS(\textbf{\textit{X}}_1,\ldots,\textbf{\textit{X}}_J)$ prior (or de Finetti measure) Q_J

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- Assess probabilities related only to observable quantities/events
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Approach (2)

- lt amounts to specifying a probability measure Q_J on an infinite-dimensional space
- It can be actually implemented

Flexible dependence structures

and discreteness

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Main findings common to all these cases:

For any A

 $\operatorname{Corr}(\tilde{p}_i(A), \tilde{p}_j(A))$ does not depend on A

and it is, then, used as an overall measure of pairwise dependence among random probabilities in $(\tilde{p}_1, \dots, \tilde{p}_J)$.

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Extreme cases

- ▶ X_i and X_j fully exchangeable \implies Corr $(\tilde{p}_i(A), \tilde{p}_j(A)) = 1$
- **X**_i and X_i unconditionally independent \implies Corr $(\tilde{p}_i(A), \tilde{p}_i(A)) = 0$

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- ► For any A

$$\operatorname{Corr}(\tilde{p}_i(A), \tilde{p}_j(A)) \geq 0$$

Open problems of interest

- ▶ Is $Corr(\tilde{p}_i(A), \tilde{p}_j(A))$ not dependent on A a general property that holds true beyond the specific examples of priors studied in the literature?
- ▶ What is the implied correlation for the observables from any two samples, i.e. $Corr(X_{i,i}, X_{\ell,i'})$?
 - ► Can $Corr(X_{i,j}, X_{\ell,j'})$ be either positive or negative?
 - What is the role played by the atoms of the underlying discrete random probability measures?
- Are there models for which
 - ightharpoonup Corr $(\tilde{p}_i(A), \tilde{p}_j(A)) = 1$ if and only if $\tilde{p}_i = \tilde{p}_j$
 - $ightharpoonup \operatorname{Corr}(\tilde{p}_i(A), \tilde{p}_i(A)) = 0$ if and only if $\tilde{p}_i \perp \tilde{p}_i$
- How is dependence reflected by the induced random partitions?
- ▶ Is it possible to provide and overall measure of dependence between \tilde{p}_i and \tilde{p}_j that captures the distance between the actual prior specification and the extremes of exchangeability and unconditional independence?
 - Ideally, an index $\mathcal{I} \in [0,1]$ such that $\mathcal{I}=0$ is equivalent to independence and $\mathcal{I}=1$ identifies the other extreme of exchangeability

Correlation with multivariate species sampling processes

Correlation structure with mSSP's

(1) If $(\mathbf{X}_1, \mathbf{X}_2)$ is a mSSS identified by (P_1, P_2) , then

$$Corr(\tilde{p}_{1}(A), \tilde{p}_{2}(A)) = \frac{\mathbb{P}[X_{1,i} = X_{2,j}]}{\sqrt{\mathbb{P}[X_{1,i} = X_{1,\ell}] \mathbb{P}[X_{2,j} = X_{2,\kappa}]}} \ge 0$$

Hence

$$\operatorname{Corr}(\tilde{p}_i(A), \tilde{p}_J(A)) = 0$$
 if and only if $\mathbb{P}[\text{"ties across samples"}] = 0$

(2) If \tilde{p}_1 and \tilde{p}_2 have the same marginal distribution, then

$$\operatorname{Corr}(\tilde{p}_1(A), \tilde{p}_2(A)) = \frac{\mathbb{P}[X_{1,i} = X_{2,j}]}{\mathbb{P}[X_{1,i} = X_{1,\ell}]} = \frac{\operatorname{Prob}[\text{"tie across samples"}]}{\operatorname{Prob}[\text{"tie within a sample"}]}$$

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- ► Corr $(\tilde{p}_i(A), \tilde{p}_i(A))$ does not depend on A for the general class of mSSP's
- ▶ $Corr(\tilde{p}_i(A), \tilde{p}_j(A))$ exclusively depends on the probabilities of sharing atoms, regardless of their specific value

$$\mathbb{P}[X_{1,i} = X_{2,j}] = \Pi_1^{(2)}(1,1)$$

$$\mathbb{P}[X_{j,i} = X_{j,\ell}] = \Phi_{1,j}^{(2)}(2) \qquad (j = 1,2).$$

Regular multivariate species sampling processes

A proper mSSP can be further rewritten as

$$\tilde{p}_{j} = \sum_{\ell=1}^{L_{0}} w_{\ell}^{(0)} \, \delta_{\theta_{\ell}} + \sum_{k=1}^{K_{j}} w_{k}^{(j)} \delta_{\eta_{j,k}}$$

- ▶ $\theta_{\ell} \stackrel{\text{iid}}{\sim} H$ are shared atoms
- $\blacktriangleright \eta_{j,k} \stackrel{\text{iid}}{\sim} H$ are sample–specific atoms
- $\qquad \qquad \blacktriangleright \ \ L_0, K_j \in \{0,1,\ldots\} \cup \{\infty\}$

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- ▶ $L_0, K_i \in \{0, 1, ...\} \cup \{\infty\}$

Regular mSSP

A mSSP $(\tilde{p}_1, \tilde{p}_2)$ is said *regular* if either

 $K_1 = K_2 = 0$ (no sample–specific atoms)

or

 $\blacktriangleright \quad (w_k^{(1)})_{k \geq 1} \bot (w_k^{(2)})_{k \geq 1} \quad \text{(independent frequencies of non--shared atoms)}$

Examples and property

- ► Hierarchical NRMIs & hierarchical Pitman–Yor processes regular mSSP since $K_1 = K_2 = 0$ Teh et al. (2006), Camerlenghi et al. (2019)
- Nested Dirichlet processes and latent nested processes regular mSSP since K₁ = K₂ = 0 Rodríguez et al. (2008), Camerlenghi et al. (2019)
- ▶ **GM–dependent processes:** regular mSSP since $K_1 = K_2 = \infty$ and $(w_k^{(1)})_{k \ge 1} \bot (w_k^{(2)})_{k \ge 1}$ Müller, Quintana and Rosner (2004), L., Nipoti and Prünster (2014)

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Property

If (P_1, P_2) is a regular mSSP, then

- $\operatorname{Corr}(\tilde{p}_1(A), \tilde{p}_2(A)) = 1$ if and only if $\tilde{p}_1 = \tilde{p}_2$ almost surely
- ► Corr($\tilde{p}_1(A), \tilde{p}_2(A)$) = 0 if and only if $\tilde{p}_1 \perp \tilde{p}_2$

Mixture representation of the pEPPF

$$\Pi_k^{(n)}(\boldsymbol{n}_1,\boldsymbol{n}_2) = \int_{\Lambda} \Pi_k^{(n)}(\boldsymbol{n}_1,\boldsymbol{n}_2;\lambda) \, \pi(\mathrm{d}\lambda)$$

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For a large family of discrete priors

- there exists $\lambda^* \in \operatorname{supp}(\pi)$ such that $\Pi_k^{(n)}(\boldsymbol{n}_1,\boldsymbol{n}_2;\lambda^*) = \Phi_k^{(n)}(\boldsymbol{n}_1+\boldsymbol{n}_2)$
- $\pi_1 = \pi(\{\lambda^*\}) > 0$

$$\Pi_k^{(N)}(\textbf{\textit{n}}_1,\textbf{\textit{n}}_2) = \pi_1 \underbrace{\Phi_k^{(N)}(\textbf{\textit{n}}_1 + \textbf{\textit{n}}_2)}_{\text{joint exchangeability}} + (1 - \pi_1) \, f_{k,N_1,N_2}(\textbf{\textit{n}}_1,\textbf{\textit{n}}_2)$$

Mixture representation of the pEPPF

$$\Pi_k^{(n)}(\boldsymbol{n}_1,\boldsymbol{n}_2) = \int_{\Lambda} \Pi_k^{(n)}(\boldsymbol{n}_1,\boldsymbol{n}_2;\lambda) \, \pi(\mathrm{d}\lambda)$$

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An example: Hidden Hierarchical Processes (L. Prünster and Rebaudo, 2022; Fasano, L., Prünster and Rebaudo, 2022+)

Hidden hierarchical Pitman-Yor processes

$$\mathsf{Nested\ component}\ \Rightarrow \left\{ \begin{array}{l} \tilde{p}_j \,|\: G \stackrel{\mathrm{iid}}{\sim} G \\ \\ G \,|\: \tilde{p}_0 \sim \mathsf{PY}(\alpha, \gamma; \mathsf{PY}(\beta, \sigma; \tilde{p}_0)) \end{array} \right.$$
 Hierarchical component $\Rightarrow \quad \tilde{p}_0 \sim Q = \mathsf{PY}(\beta_0, \sigma_0; H)$

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Definition

A vector $(\tilde{p}_1, \dots, \tilde{p}_J)$ is a hidden hierarchical Pitman–Yor process (HHPY) if

$$\begin{split} \tilde{p}_{j}|G &\stackrel{\text{iid}}{\sim} G = \sum_{k \geq 1} \omega_{k} \, \delta_{P_{k}^{*}} & (\omega_{k})_{k \geq 1} \sim \mathsf{GEM}(\alpha, \gamma) \\ \\ P_{k}^{*}|\tilde{p}_{0} &\stackrel{\text{iid}}{\sim} \mathsf{PY}(\beta, \sigma; \tilde{p}_{0}) & \tilde{p}_{0} \sim \mathsf{PY}(\beta_{0}, \sigma_{0}; H) \end{split}$$

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Notation: $(\tilde{p}_1, \dots, \tilde{p}_J) \sim \mathsf{HHPY}(\psi; P_0)$, where $\psi = (\alpha, \gamma, \sigma, \beta, \sigma_0, \beta_0)$

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 $\gamma = \sigma = \sigma_0 = 0$ \Longrightarrow Hidden hierarchical Dirichlet process (HHDP)

pEPPF

When
$$J=2$$

$$\Pi_k^{(N)}(\textbf{\textit{n}}_1,\textbf{\textit{n}}_2) = \frac{1-\gamma}{\alpha+1} \Phi_{k,1}^{(N)}(\textbf{\textit{n}}_1+\textbf{\textit{n}}_2) + \frac{\alpha+\gamma}{\alpha+1} \Phi_{k,2}^{(N)}(\textbf{\textit{n}}_1,\textbf{\textit{n}}_2)$$

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With $\mathscr{C}(n, k; \sigma)$ denoting the generalized factorial coefficient

$$\begin{aligned} \Phi_{k,1}^{(N)}(\mathbf{n}_1 + \mathbf{n}_2) &= \frac{\prod_{r=1}^{k-1} (\beta_0 + r\sigma_0)}{(\beta + 1)_{N-1}} \sum_{\ell} \frac{\prod_{s=1}^{|\ell|-1} (\beta + s\sigma)}{(\beta_0 + 1)_{|\ell|-1}} \\ &\times \prod_{j=1}^{k} \frac{\mathscr{C}(\mathbf{n}_{1,j} + \mathbf{n}_{2,j}, \ell_j; \sigma)}{\sigma^{\ell_j}} (1 - \sigma_0)_{\ell_j - 1} \end{aligned}$$

$$\Phi_{k,2}^{(N)}(\pmb{n}_1,\pmb{n}_2)=$$
 available, though it does not correspond to independence $eq \Phi_{k,1}^{(N_1)}(\pmb{n}_1)\Phi_{k,2}^{(N_2)}(\pmb{n}_2)$

Posterior probability of exchangeability

$$\mathbb{P}[\tilde{p}_1 = \tilde{p}_2 \,|\, \textbf{\textit{X}}_1, \textbf{\textit{X}}_2] = \frac{(1 - \gamma) \, \Phi_{k,1}^{(N)}(\textbf{\textit{n}}_1 + \textbf{\textit{n}}_2)}{(1 - \gamma) \Phi_{k,1}^{(N)}(\textbf{\textit{n}}_1 + \textbf{\textit{n}}_2) + (\alpha + \gamma) \Phi_{k,2}^{(N)}(\textbf{\textit{n}}_1, \textbf{\textit{n}}_2)}$$

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Number of clusters K_N

▶ Distribution of the number of clusters K_N out of $N = N_1 + N_2$ observations from the two samples

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- ▶ Distribution of the number of clusters K_N out of $N = N_1 + N_2$ observations from the two samples
- ► Asymptotics of K_N

▶ If
$$N_1 = N_2 = N/2$$
 and $\sigma, \sigma_0 \in (0, 1)$, as $N \to \infty$

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If
$$N_1 = N_2 = N/2$$
 and $\sigma = \sigma_0 = 0$, as $N \to \infty$

$$K_N \simeq \log \log N$$
 a.s.

Dependence and random atoms: beyond mSSP's

If one sticks to a mSSP for partially exchangeable data ($\textbf{X}_1, \textbf{X}_2$), then

- (1) $\operatorname{Corr}(\tilde{\rho}_1(A), \tilde{\rho}_2(A)) \geq 0$
- (2) $\operatorname{Corr}(X_{1,\ell}, X_{2,\kappa}) \geq 0$

Dependence and random atoms: beyond mSSP's

If one sticks to a mSSP for partially exchangeable data (X_1, X_2) , then

- (1) $Corr(\tilde{p}_1(A), \tilde{p}_2(A)) > 0$
- (2) $Corr(X_1, X_2, X_2) > 0$

How recover a wider range of correlations?

▶ Bivariate SSP on X²

$$p_1^* = \sum_{h \ge 1} w_h^{(1)} \, \delta_{(\theta_h, \phi_h)}, \quad p_2^* = \sum_{h \ge 1} w_h^{(2)} \, \delta_{(\theta_h, \phi_h)}, \quad (\theta_h, \phi_h) \stackrel{\text{iid}}{\sim} G_0$$

Projections

$$\boxed{\tilde{p}_1(\,\cdot\,) = p_1^*(\,\cdot\,\times\mathbb{X}) = \sum_{h \geq 1} w_h^{(1)}\,\delta_{\theta_h}} \boxed{\tilde{p}_2(\,\cdot\,) = p_2^*(\mathbb{X}\times\,\cdot\,) = \sum_{h \geq 1} w_h^{(2)}\,\delta_{\phi_h}}$$

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$$(\tilde{p}_1, \tilde{p}_2)$$
 is a bivariate SSP iff: **(1)** $G_0 = P_0^2$ (independence) or **(2)** $G_0(\{\theta_h = \phi_h, \text{ for any } h\}) = 1$ (shared atoms)

Correlations with any sign

Probability of a tie within sample 1 or sample 2

$$\beta = \mathbb{P}[X_{1,i} = X_{1,j}] = \mathbb{P}[X_{2,i'} = X_{2,j'}] = \sum_{k \ge 1} \mathbb{E}\{w_k^{(1)}\}^2$$

Probability of a hyper–tie between sample 1 and sample 2

$$\gamma = \mathbb{P}[(X_{1,1}, X_{2,1}) = (\theta_k, \phi_k) \text{ for some } k] = \sum_{k \ge 1} \mathbb{E} w_k^{(1)} w_k^{(2)}$$

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Ties vs hyper-ties

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Ties vs hyper-ties

- ▶ $0 \le \gamma \le \beta$ ▶ $\gamma = \beta$ if and only if $w_k^{(1)} \stackrel{\text{a.s.}}{=} w_k^{(2)}$, for any k.

How do the probabilities of tie (β) and of hyper–tie (γ) relate to correlations within and between samples?

Ties and hyper-ties probabilities

Correlations and ties/hyper-ties

$$\mathsf{Corr}(X_{1,i},X_{1,j}) = \mathsf{Corr}(X_{2,i'},X_{2,j'}) = \beta$$

$$\operatorname{Corr}(X_{1,1}, X_{2,1}) = \gamma \rho_0$$
 where $\rho_0 = \operatorname{Corr}(\theta_1, \phi_1)$

Ties and hyper-ties probabilities

Correlations and ties/hyper-ties

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Corollary

- **1)** $Corr(X_{1,1}, X_{2,1}) \in [-\beta, \beta]$
- **2)** $|\text{Corr}(X_{1,1}, X_{2,1})| = \beta$ if and only if: **2.1)** $\omega_k^{(1)} \stackrel{\text{a.s.}}{=} \omega_k^{(2)}$ for any k **2.2)** $|\rho_0| = 1$

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Full Range of Borrowing Information (FuRBI): independence shrinkage repulsion / negative correlation

An effective proposal: nFuRBI priors

Completely random vector

$$(\tilde{\mu}_1, \tilde{\mu}_2) = \Big(\sum_{h \geq 1} J_h \, \delta_{(\theta_h, \phi_h)}, \sum_{h \geq 1} W_h \, \delta_{(\theta_h, \phi_h)}\Big)$$

where $(J_h, W_h, \theta_h, \phi_h)_{h>1}$ are points of a Poisson process with Lévy intensity

$$\nu(\mathrm{d}s_1,\mathrm{d}s_2,\mathrm{d}\theta,\mathrm{d}\phi) = \rho(s_1,s_2)\mathrm{d}s_1\mathrm{d}s_2\ G_0(\mathrm{d}\theta,\mathrm{d}\phi)$$

and
$$G_0(\cdot \times \mathbb{X}) = G_0(\mathbb{X} \times \cdot) = P_0(\cdot)$$

▶ Projections of $(\tilde{\mu}_1, \tilde{\mu}_2)$

$$\mu_1(\,\cdot\,) = \tilde{\mu}_1(\,\cdot\,\times\,\mathbb{X}), \quad \mu_2(\,\cdot\,) = \tilde{\mu}_2(\mathbb{X}\times\,\cdot\,)$$

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$$\mu_1(\cdot) = \tilde{\mu}_1(\cdot \times \mathbb{X}), \quad \mu_2(\cdot) = \tilde{\mu}_2(\mathbb{X} \times \cdot)$$

nFuRBI priors

If ho is such that $\int_{\mathbb{R}^2_+}
ho(s_1,s_2) \, \mathrm{d} s_1, \mathrm{d} s_2 = \infty$, then

$$(\tilde{p}_1, \tilde{p}_2) = (\mu_1/\mu_1(\mathbb{X}), \, \mu_2/\mu_2(\mathbb{X}))$$

is a normalized FuRBI CRM or, in short, nFuRBI prior.

With $k_i \leq N_1$ and $k_2 \leq N_2$, let

- (1) $X_{1,1}^*, \dots, X_{1,k_1}^*$ distinct values in $X_1^{(N_1)}$
- (2) $X_{2,1}^*, \dots, X_{2,k_2}^*$ distinct values in $\mathbf{X}_2^{(N_2)}$

Predictive distribution

Conditional on samples $\boldsymbol{X}_1^{(N_1)}$ and $\boldsymbol{X}_2^{(N_2)}$ as in (1)–(2), one has

$$X_{1,N_{1}+1} | (\boldsymbol{X}_{1}^{(N_{1})}, \boldsymbol{X}_{2}^{(N_{2})}) \sim \xi_{0} P_{0} + \sum_{i=1}^{k_{1}} \xi_{1,i} \delta_{X_{1,i}^{*}} + \sum_{i=1}^{k_{2}} \xi_{2,i} P_{X_{2,i}^{*}}$$

$$X_{2,N_2+1} | (\boldsymbol{X}_1^{(N_1)}, \boldsymbol{X}_2^{(N_2)}) \sim \eta_0 P_0 + \sum_{i=1}^{k_2} \eta_{2,i} \, \delta_{X_{2,i}^*} + \sum_{i=1}^{k_1} \eta_{1,i} P_{X_{1,i}^*}$$

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Conditional on samples $\mathbf{X}_1^{(N_1)}$ and $\mathbf{X}_2^{(N_2)}$ as in (1)–(2), one has

$$X_{1,N_1+1} \mid (\boldsymbol{X}_1^{(N_1)}, \boldsymbol{X}_2^{(N_2)}) \sim \xi_0 P_0 + \sum_{i=1}^{k_1} \xi_{1,i} \delta_{X_{1,i}^*} + \sum_{i=1}^{k_2} \xi_{2,i} P_{X_{2,i}^*}$$

$$X_{2,N_2+1} | (\boldsymbol{X}_1^{(N_1)}, \boldsymbol{X}_2^{(N_2)}) \sim \eta_0 P_0 + \sum_{i=1}^{k_2} \eta_{2,i} \, \delta_{X_{2,i}^*} + \sum_{i=1}^{k_1} \eta_{1,i} P_{X_{1,i}^*}$$

Posterior distribution

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If $\mathbf{X}_1^{(N_1)}$ and $\mathbf{X}_2^{(N_2)}$ are samples as in (1)–(2), conditional on a suitable set of latent variables, the posterior distribution of P_1 equals the distribution of the random probability measure

$$w_{1,n}P_1^* + w_{2,n} \sum_{i} \pi_{1,i} \, \delta_{X_{1,i}^*} + (1 - w_{2,n} - w_{3,n}) \sum_{j} \pi_{2,j} \delta_{Z_{X_{2,j}^*}}$$

where $Z_{X_{2,j}^*} \stackrel{\text{ind}}{\sim} P_{X_{2,j}^*}$. Similar representation holds true for P_2 .

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Mixture of

- ► an updated nFURBI (P₁*)
- a weighted empirical distribution of the sample data
- ▶ a weighted empirical distribution of a transformation of the other sample's data $\Rightarrow G_0\{\theta_k = \phi_k, \text{ for any } k\} = 1, \text{ then } Z_{X_{2,j}^*} = X_{2,j}^*$: standard posterior characterization of random measure—beased models.
- ► Talk by Beatrice Franzolini (Monday)

Measuring dependence

Several priors for partially exchangeable data are transformations of completely random vectors $\tilde{\mu} = (\tilde{\mu}_1, \dots, \tilde{\mu}_J)$, i.e.

$$\tilde{p}_j = T(\tilde{\mu}_j), \qquad i = 1, \dots, J$$

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- ► Two extreme situations
 - lacktriangle Complete dependence or co-monotonicity $ilde{\mu}^{ extsf{CO}}$

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A possible index

For some distance d and completely random vector $ilde{oldsymbol{\mu}}$

$$\mathcal{I}_{d}(\tilde{\boldsymbol{\mu}}) = 1 - \frac{d^{2}(\tilde{\boldsymbol{\mu}}, \tilde{\boldsymbol{\mu}}^{CO})}{\sup_{\boldsymbol{\mu}'} d^{2}(\boldsymbol{\mu}', \tilde{\boldsymbol{\mu}}^{CO})}$$

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Which d? Based on an extended Wasserstein distance d_W between the corresponding Lévy intensities ν^1 and ν^2

$$\mathcal{W}(\tilde{\mu}_1, \tilde{\mu}_2) = \sup_A d_W(\nu_A^1, \nu_A^2)$$

The Wasserstein based index of dependence

If for some completely random vector $\tilde{\mu}$, a prior is defined through $\tilde{p}_i = T(\tilde{\mu}_i)$, how close is such a specification from the exchangeability extreme?

For any completely random vector $ilde{m{\mu}}, \mathcal{I}_{\mathcal{W}}(ilde{m{\mu}}) \in [0,1].$ Moreover

- $lacksquare \mathcal{I}_{\mathcal{W}}(ilde{\mu}) = 1$ if and only if $ilde{\mu} = ilde{\mu}^{\mathsf{CO}}$
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- ► Evaluation for some classes of priors
 - $\tilde{\mu}$ a completely random vector with an additive structure (L., Nipoti and Prünster, 2014)
 - Compound random measures (Griffin and Leisen, 2017)
- Possible extension to posterior distribution would allow testing for distributional homogeneity

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Talks by Marta Catalano (Wednesday) and Hugo Lavenant (Tuesday)

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