SURVIVAL ANALYSIS VIA HIERARCHICALLY DEPENDENT MIXTURE HAZARDS

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Hierarchical nonparametric processes are popular tools for defining priors on collections of probability distributions, which induce dependence across multiple samples. In survival analysis problems, one is typically interested in modeling the hazard rates, rather than the probability distributions themselves, and the currently available methodologies are not applicable. Here, we fill this gap by introducing a novel, and analytically tractable, class of multivariate mixtures whose distribution acts as a prior for the vector of sample-specific baseline hazard rates. The dependence is induced through a hierarchical specification of the mixing random measures that ultimately corresponds to a composition of random discrete combinatorial structures. Our theoretical results allow to develop a full Bayesian analysis for this class of models, which can also account for right-censored survival data and covariates, and we also show posterior consistency. In particular, we emphasize that the posterior characterization we achieve is the key for devising both marginal and conditional algorithms for evaluating Bayesian inferences of interest. The effectiveness of our proposal is illustrated through some synthetic and real data examples.

1. Introduction. Hierarchical processes are hugely popular Bayesian nonparametric models, which have seen successful applications in linguistics, information retrieval, topic modeling and genomics, among others. They are ideally suited to model relationships across multiple samples, which may share distinct observations' values (or latent features, if used in a mixture setup). For instance, a topic might be shared by different documents of a *corpus* or a specific subsequence of a cDNA sequence might be recorded at different tissues of an organism. The prototype of this class of models is the *hierarchical Dirichlet process* (HDP) introduced in [41], which can be seen as an infinite-dimensional extension of the latent Dirichlet allocation model in [3]. Besides being a clever construction, the availability of suitable algorithms allow to perform Bayesian inference. The general structure of hierarchical processes amounts to

$$(1) \quad \begin{aligned} (\tilde{p}_1, \dots, \tilde{p}_d) | \tilde{p}_0 &\overset{\text{i.i.d.}}{\sim} \tilde{\mathcal{L}}_0, \\ \tilde{p}_0 &\sim \mathcal{L}_0, \end{aligned}$$

where $\tilde{\mathcal{L}}_0$ is the probability distribution of each random probability measure $\tilde{p}_i$ such that $\mathbb{E}[\tilde{p}_i | \tilde{p}_0] = \int p \tilde{\mathcal{L}}_0(\mathrm{d}p) = \tilde{p}_0$, and $\mathcal{L}_0$ is such that $\mathbb{E}[\tilde{p}_0] = \int p \mathcal{L}_0(\mathrm{d}p) = P_0$, for some fixed nonatomic probability measure P_0. The vector of random probability measures in (1) defines a prior for the probability distributions of d partially exchangeable samples with dependence across samples being induced by $\tilde{p}_0$. Since $\tilde{p}_0$ and $(\tilde{p}_1, \dots, \tilde{p}_d)$ are taken to be discrete random probability measures, distinct values are shared within and across the d samples.

Clearly, in the extreme case of $\tilde{p}_0$ degenerating on P_0, the $\tilde{p}_i$'s are unconditionally independent corresponding to independence across samples with no shared values; see [40, 41] and, for extensions to general random measures and a complete picture of their distributional properties, [4].

Though currently available hierarchical processes are effective in modeling probability measures, in survival analysis one needs a flexible tool for modeling directly the hazard rates. Here we successfully address the issue and introduce the natural counterpart, for hazard rate functions, of hierarchical random probability measures. This corresponds to a class of dependent hazard models defined as mixtures with mixing measures displaying a hierarchical dependence structure. Such a prior is ideally suited to model hazard rate functions associated to different, though related, populations in a similar fashion as hierarchical processes do for related probability distributions. Moreover, it is possible to incorporate both censored observations and covariates in a straightforward way. More specifically, our proposal assumes a prior for the hazard rate functions of each individual sample identified by the distribution of the random mixture

$$(2) \qquad \tilde{h}_\ell(t) = \int_{\mathbb{Y}} k(t; y) \tilde{\mu}_\ell(\mathrm{d}y), \quad \ell = 1, \ldots, d,$$

where $k(\cdot; \cdot)$ is a suitable kernel and $\tilde{\mu}_\ell$ a random measure. Note that an extension of such a proposal that accommodates also for subject-specific covariates can be achieved by resorting, for example, to a semiparametric representation through a multiplicative term as in Cox proportional hazards models. Dependence among the hazard rates is then created at the level of the random measures through a hierarchical structure

$$(3) \qquad \begin{aligned} (\tilde{\mu}_1 \ldots \tilde{\mu}_d)|\tilde{\mu}_0 &\overset{\text{i.i.d.}}{\sim} \tilde{\mathcal{G}}_0, \\ \tilde{\mu}_0 &\sim \mathcal{G}_0, \end{aligned}$$

where $\tilde{\mathcal{G}}_0$ is the distribution of each random measure $\tilde{\mu}_\ell$ and depends on $\tilde{\mu}_0$, which in turn is distributed according to $\mathcal{G}_0$. An important feature of the proposed construction is that it allows for nonproportional hazard rates across samples. In the sequel $\tilde{\mu}_0$ and $\tilde{\mu}_\ell|\tilde{\mu}_0$, for $\ell = 1, \ldots, d$, will be assumed to be completely random measures (CRMs), which play a key role in Bayesian nonparametrics as effectively described in [29]. This will allow, in the following sections, to derive some key distributional properties of $(\tilde{\mu}_0, \tilde{\mu}_1, \ldots, \tilde{\mu}_d)$, both *a priori* and *a posteriori*. Based on the latter, and conditional on a suitable latent structure, closed-form expressions for posterior estimates of both the hazard rates and the survival functions are obtained as well as a sampling scheme for performing full Bayesian inference.

Survival analysis has been one of the driving application areas for Bayesian nonparametrics since its early days and among the many seminal papers in the simple exchangeable setup we mention [12, 15, 17, 19, 22, 30]. The latter two papers are of particular importance to the present contribution since they introduced the mixture hazard structure (2) with a gamma process as mixing measure. Still in the exchangeable case, they were extended to general mixing CRMs in [25] and further investigated in, for example, [9, 13, 23, 35–38]. Beyond the exchangeable case, a popular modeling strategy, which yields the desired heterogeneity across samples, relies on the construction of dependent and sample-specific random measures that, suitably transformed, allow to model partially exchangeable data. An example is the mixture transformation in (10), which is the focus of this contribution. This technique has been widely used for creating dependent random *probability* measures in the literature and examples are available in, for example, [5, 10, 11, 16, 28, 32, 34]. For a review, see [33]. The only contribution aiming to model directly dependent hazard rates, is [27], which however has the drawback of incurring into combinatorial, and hence computational, issues for $d > 2$ samples. In contrast, the methodology proposed here leads to simple and tractable expressions for $d > 2$ groups of observations.

The outline of the paper is as follows. In Section 2, the basic building blocks of our model, namely vectors of hierarchical CRMs, are defined. Hierarchically dependent random hazard rates are introduced and investigated in Section 3. Posterior consistency, in a partially exchangeable framework, is established in Section 4, while the posterior distribution is identified in Section 5. The theoretical findings form the basis for devising suitable marginal and conditional sampling schemes, which are described in Section 6. Finally, a simulation study is considered in Section 7. Proofs, other relevant technical details, extensions accounting for right-censored data and covariates as well as additional illustrations with real data are deferred to the Supplementary Material [6].

2. Hierarchical random measures. The key ingredient for our dependent hazard rates model is a vector of random measures $(\tilde{\mu}_1, \ldots, \tilde{\mu}_d)$ with dependence induced by a hierarchical structure as in (3). First, we introduce the notion of completely random measure (CRM). A CRM is a random element $\tilde{\mu}$, defined on some probability space $(\Omega, \mathscr{F}, \mathbb{P})$ and taking values in the space $\mathsf{M}_\mathbb{Y}$ of boundedly finite measures on some Polish space $\mathbb{Y}$, such that $\tilde{\mu}(A_1), \ldots, \tilde{\mu}(A_n)$ are mutually independent random variables for any choice of bounded and pairwise disjoint Borel sets $A_1, \ldots, A_n$, and any $n \geq 1$. In the following, we consider CRMs without fixed jumps and no deterministic component. They are characterized by the respective Laplace functional transform at any measurable function $g : \mathbb{Y} \to \mathbb{R}^+$,

$$(4) \qquad \mathbb{E}\big[e^{-\int_\mathbb{Y} g(y)\tilde{\mu}(\mathrm{d}y)}\big] = \exp\bigg\{-\int_{\mathbb{R}^+ \times \mathbb{Y}} (1 - e^{-sg(y)})\nu(\mathrm{d}s, \mathrm{d}y)\bigg\},$$

where ν, known as the Lévy intensity measure, uniquely identifies $\tilde{\mu}$. Hence, the notation $\tilde{\mu} \sim \mathrm{CRM}(\nu)$. For completeness, we remind that $\tilde{\mu}$ may be seen as a functional of a Poisson random measure $\tilde{N} = \sum_{i \geq 1} \delta_{(J_i, Y_i)}$ on $\mathbb{R}^+ \times \mathbb{Y}$ characterized by a mean intensity measure ν such that for any Borel set A in $\mathbb{R}^+ \times \mathbb{Y}$ with $\nu(A) < \infty$ one has $\tilde{N}(A) \sim \mathrm{Po}(\nu(A))$. The CRM $\tilde{\mu}$ can be then represented as $\sum_{i \geq 1} J_i \delta_{Y_i}$, hence its realizations are a.s. discrete with both jumps and locations random. See [26] for an exhaustive account on CRMs.

In analogy to the hierarchical construction for random probability measures, we define a hierarchical structure for CRMs as

$$(5) \qquad \begin{aligned} \tilde{\mu}_\ell | \tilde{\mu}_0 &\overset{\mathrm{ind}}{\sim} \mathrm{CRM}(\tilde{\nu}_\ell), \quad \ell = 1, \ldots, d \\ \tilde{\mu}_0 &\sim \mathrm{CRM}(\nu_0), \end{aligned}$$

with Lévy intensities

$$(6) \qquad \tilde{\nu}_\ell(\mathrm{d}s, \mathrm{d}y) = \rho_\ell(s)\,\mathrm{d}s\,\tilde{\mu}_0(\mathrm{d}y), \qquad \nu_0(\mathrm{d}s, \mathrm{d}y) = \rho_0(s)\,\mathrm{d}s\,c_0 P_0(\mathrm{d}y),$$

where P_0 is a diffuse probability measure on $\mathbb{Y}$, while ρ_ℓ (for $\ell = 1, \ldots, d$) and ρ_0 are nonnegative measurable functions such that $\int_0^\infty \min\{1, s\}\rho_\ell(s)\,\mathrm{d}s < \infty$ and $\int_0^\infty \min\{1, s\} \times \rho_0(s)\,\mathrm{d}s < \infty$.

REMARK 1. The specification in (5) entails that the $\tilde{\mu}_\ell$'s may have different distributions, which yields greater flexibility in applications. Nonetheless, for the sake of simplicity we henceforth stick to the case where $\rho_\ell = \rho$ for any $\ell = 1, \ldots, d$. The results we obtain can be adapted to recover the case where the ρ_ℓ's differ and we defer this to future work.

The fundamental tool to work with is the Laplace functional transform of a hierarchical CRM vector $(\tilde{\mu}_1, \ldots, \tilde{\mu}_d)$, which has a simple structure. For $\ell = 1, \ldots, d$, set $\tilde{\mu}_\ell(g_\ell) = \int_\mathbb{Y} f_\ell(y)\tilde{\mu}_\ell(\mathrm{d}y)$ with g_ℓ a nonnegative real valued function. Moreover, we let $\psi^{(0)}(u) :=$

$\int_0^\infty (1 - e^{-su})\rho_0(s)\,\mathrm{d}s$ and $\psi^{(\ell)}(u) := \int_0^\infty (1 - e^{-su})\rho_\ell(s)\,\mathrm{d}s$, as $\ell = 1, \ldots, d$. By exploiting conditional independence and (4), the Laplace functional transform is given by

$$(7) \qquad \mathbb{E}\big[e^{-\tilde{\mu}_1(g_1) - \cdots - \tilde{\mu}_d(g_d)}\big] = \exp\bigg\{-c_0 \int_{\mathbb{Y}} \psi^{(0)}\bigg[\sum_{\ell=1}^d \psi^{(\ell)}(g_\ell(y))\bigg] P_0(\mathrm{d}y)\bigg\}.$$

Now we introduce two special cases to be considered throughout and which represent natural choices for hierarchical CRM constructions.

EXAMPLE 1 (Hierarchical gamma CRM model). If both $\tilde{\mu}_\ell$'s and $\tilde{\mu}_0$ are gamma CRMs, which corresponds to $\rho_0(s) = \rho_\ell(s) = e^{-s}s^{-1}$ with $c_0 = 1$, we obtain the hierarchical gamma CRM model, whose Laplace functional reduces to

$$\mathbb{E}\big[e^{-\tilde{\mu}_1(g_1) - \cdots - \tilde{\mu}_d(g_d)}\big] = \exp\bigg\{-\int_{\mathbb{Y}} \log\Big(1 + \sum_{\ell=1}^d \log\big(1 + g_\ell(y)\big) P_0(\mathrm{d}y)\bigg\}.$$

EXAMPLE 2 (Hierarchical generalized gamma CRM model). Let us assume that both $\tilde{\mu}_\ell$'s and $\tilde{\mu}_0$ are generalized gamma CRMs, that is, $\rho_0(s) = e^{-s}s^{-1-\sigma_0}/\Gamma(1-\sigma_0)$ and $\rho_\ell(s) = e^{-s}s^{-1-\sigma}/\Gamma(1-\sigma)$ for some σ and σ_0 in $(0, 1)$, we obtain the hierarchical generalized gamma CRM model with Laplace functional

$$\mathbb{E}\big[e^{-\tilde{\mu}_1(g_1) - \cdots - \tilde{\mu}_d(g_d)}\big]$$

$$= \exp\bigg\{-\frac{c_0}{\sigma_0} \int_{\mathbb{Y}} \bigg[\bigg(\sum_{\ell=1}^d \frac{(g_\ell(y) + 1)^\sigma - 1}{\sigma} + 1\bigg)^{\sigma_0} - 1\bigg] P_0(\mathrm{d}y)\bigg\}.$$

REMARK 2. The construction of hierarchical CRMs in (5) is closely related to the popular Bochner subordination in the theory of Lévy processes. Indeed, with $\mathbb{Y} = [0, 1]$, define $t \mapsto \tau_t = \tilde{\mu}_0((0, t])$ and $t \mapsto \xi_t^{(\ell)} = \tilde{\mu}_\ell((0, t])$. Moreover, let $P_0(\mathrm{d}t) = \mathrm{d}t$. In this case, $\{\tau_t : 0 \leq t \leq 1\}$ is known as a *subordinator* and the time-changed process $\{\xi_{\tau_t}^{(\ell)} : 0 \leq t \leq 1\}$ corresponds to the so-called Bochner's subordination; see, for example, [2] and [39]. Hence, our proposal can be seen as an extension of such a construction to more abstract spaces. Subordination of Lévy processes has been widely used in Finance and the first contributions in this direction can be found in [31] and [7]. Very much in the spirit of the present paper, subordination has been applied to a vector of independent Brownian motions $(B_{\tau_t}^{(1)}, \ldots, B_{\tau_t}^{(d)})$ in order to define dependent Lévy processes for financial applications. Examples can be found in the monograph by Cont and Tankov [8].

3. Hierarchical mixture hazard rates. If an exchangeable sequence of lifetimes, or time-to-event data, is from an absolutely continuous distribution, with hazard rate h, the pioneering papers by [15] and [30] define a prior for h in terms of a mixture model. Recast in a more general setting, as done in [25], their prior equals the distribution of the random hazard

$$(8) \qquad t \mapsto \tilde{h}(t) := \int_{\mathbb{Y}} k(t; y)\tilde{\mu}(\mathrm{d}y),$$

where $\tilde{\mu}$ is a mixing CRM and $k : \mathbb{R}^+ \times \mathbb{Y} \to \mathbb{R}^+$ is a suitable transition kernel. Dykstra and Laud [15] considered the case of $\tilde{h}$ being an *extended gamma process*, which corresponds to $\tilde{\mu}$ being a gamma CRM and $k(t; y) = \mathbb{1}_{(0,t]}(y)\alpha(y)$ for some positive and right-continuous function α. Lo and Weng [30] investigated the *weighted gamma process*, which arises with $\tilde{\mu}$ still a gamma CRM and k a general kernel. The posterior characterization for the general case was first derived in [25].

Assuming the lifetimes are exchangeable is clearly restrictive for practical purposes. A typical example may refer to data collected under d different experimental conditions that identify d samples or groups such as patients suffering from the same illness but undergoing different treatments resulting in homogeneity within the groups of patients undergoing the same treatment and heterogeneity across groups. It is worth noting that the d samples may be identified also by a vector of categorical covariates, as long as they take on finitely many values. For example, the real data illustration in the Supplementary Material (Section S7) discusses the case where the patients' groups are determined by the hospital where they are treated and by the type of tumor they suffer from. In these cases, partial exchangeability [20] is the appropriate notion of dependence, since it implies that observations are exchangeable within the same group and conditionally independent across the different groups. More precisely, for an array $\{(X_{\ell,i})_{i\geq 1} : \ell = 1, \ldots, d\}$ of $\mathbb{X}$-valued random elements we assume that for any $\ell_1 \neq \cdots \neq \ell_k$ in $\{1, \ldots, d\}$, $(i_1, \ldots, i_k) \in \mathbb{N}^k$ and $k \geq 1$,

$$(9) \qquad (X_{\ell_1,i_1}, \ldots, X_{\ell_k,i_k})|(\tilde{p}_1, \ldots, \tilde{p}_d) \overset{\text{i.i.d.}}{\sim} \tilde{p}_{\ell_1} \times \cdots \times \tilde{p}_{\ell_k},$$
$$(\tilde{p}_1, \ldots, \tilde{p}_d) \sim Q,$$

with Q is a distribution on $\mathsf{P}_{\mathbb{X}}^d$ dictating the dependence across the different groups. The literature on dependent nonparametric priors essentially boils down to the definition of probability measures Q inducing natural dependence structures across groups while still preserving mathematical tractability. Hierarchical processes represent one of the most popular instances.

In the survival context, that is, $\mathbb{X} = \mathbb{R}^+$, it is often more convenient to devise inferential procedures that are suited for models based on hazard rates and this motivates our approach; in addition our proposal allows for a straightforward inclusion of possible censoring mechanisms and individuals' covariates as discussed in the Supplementary Material (see Section S7). The prior specification we suggest resorts to a vector of dependent hazard rates $\tilde{h}_1, \ldots, \tilde{h}_d$ based on hierarchically dependent CRMs according to the following.

DEFINITION 1. A partially exchangeable sequence $\{(X_{\ell,i})_{i\geq 1} : \ell = 1, \ldots, d\}$ is directed by a *hierarchically dependent mixture hazard model* if it is characterized by (9) with

$$(10) \qquad \tilde{p}_\ell((-\infty, t]) = \mathbb{1}_{(0,+\infty)}(t) \exp\left(-\int_0^t \tilde{h}_\ell(s)\,\mathrm{d}s\right) \quad (\ell = 1, \ldots, d),$$

where the hazard rates $\tilde{h}_\ell$ admit the mixture representation in (2) and the mixing hierarchical CRMs are defined as in (3).

Note that our model allows for nonproportional hazards across samples. Moreover, the survival function associated to the hazard rate (10) is

$$(11) \qquad \tilde{S}_\ell(t) = \exp\left(-\int_{\mathbb{Y}} K_t^{(\ell)}(y)\tilde{\mu}_\ell(\mathrm{d}y)\right), \quad \ell = 1, \ldots, d,$$

where $K_t^{(\ell)}(y) = \int_0^t k(s; y)\,\mathrm{d}s$ for each $\ell = 1, \ldots, d$. Two important properties of the dependent survival functions are the determination of sufficient conditions guaranteeing they are proper and of their pairwise correlation structure. These are provided in the following result.

THEOREM 1. *Consider a hierarchical mixture hazard model as in Definition* 1.

(i) *If $\int_0^\infty \rho_\ell(s)\,\mathrm{d}s = +\infty$, for $\ell = 0, \ldots, d$, and $\lim_{t\to\infty} \int_0^t k(s; y)\,\mathrm{d}s = \infty$, P_0-a.s., then with probability* 1

$$(12) \qquad \lim_{t\to\infty} \tilde{S}_\ell(t) = 0 \quad (\ell = 1, \ldots, d).$$

(ii) *The covariance between any two survival functions $\ell_1 \neq \ell_2$, time points $t_1, t_2 \in \mathbb{R}^+$ equals*

$$\text{Cov}(\tilde{S}_{\ell_1}(t_1), \tilde{S}_{\ell_2}(t_2))$$

(13)
$$= \exp\left(-c_0 \int_{\mathbb{Y}} \psi^{(0)}\left(\sum_{i=1}^{2} \psi^{(\ell_i)}(K_{t_i}^{(\ell_i)}(y))\right) P_0(dy)\right)$$

$$- \exp\left(-c_0 \sum_{i=1}^{2} \int_{\mathbb{Y}} \psi^{(0)}(\psi^{(\ell_i)}(K_{t_i}^{(\ell_i)}(y)) P_0(dy)\right) \geq 0.$$

As for (ii), the nonnegativity of the covariance between random probabilities $\tilde{p}_i(A)$ and $\tilde{p}_j(A)$, for any $i \neq j$, is a common feature of a number of dependent processes' proposals in the literature. In particular, there are several instances where $\text{Cov}(\tilde{p}_i(A), \tilde{p}_j(A))$ does not even depend on the specific set A and is then often interpreted as a measure of overall dependence between $\tilde{p}_i$ and $\tilde{p}_j$.

Theorem 1 can be nicely illustrated in two special cases that correspond to the hierarchical CRMs already considered in Section 2. Note further that we shall consider the $\tilde{\mu}_\ell$'s conditionally identically distributed, that is, $\rho_\ell = \rho$ for any ℓ, and hence it is not surprising that in both examples $\text{Cov}(\tilde{S}_{\ell_1}(t_1), \tilde{S}_{\ell_2}(t_1))$ does not depend on the specific $\ell_1 \neq \ell_2$.

EXAMPLE 3 (Hierarchical gamma mixture hazard model). Consider the Dykstra–Laud kernel $k(t; y) := \mathbb{1}_{(0,t]}(y)\alpha(y)$ and, for simplicity, take $\alpha(y) \equiv \alpha > 0$. Combining this kernel with a hierarchical vector of gamma CRMs on $\mathbb{Y} = \mathbb{R}^+$ leads to a hierarchical gamma mixture hazard model. The associated survival functions are proper, given they clearly satisfy the conditions of (i) in Theorem 1. As for the pairwise correlation structure, by (ii) in Theorem 1 and calculations detailed in the Supplementary Material, one obtains for any $\ell_1 \neq \ell_2$ and $t_1 \leq t_2$ in $\mathbb{R}^+$,

$$\text{Cov}(\tilde{S}_{\ell_1}(t_1), \tilde{S}_{\ell_2}(t_2))$$

$$= \exp\left(-c_0 \int_{t_1}^{t_2} \log(1 + \log(1 + \alpha(t_2 - y))) P_0(dy)\right)$$

(14)
$$\times \left\{\exp\left(-c_0 \int_{0}^{t_1} \log\left(1 + \sum_{i=1}^{2} \log(1 + \alpha(t_i - y))\right) P_0(dy)\right)\right.$$

$$\left. - \exp\left(-\sum_{i=1}^{2} c_0 \int_{0}^{t_1} \log(1 + \log(1 + \alpha(t_i - y))) P_0(dy)\right)\right\}.$$

EXAMPLE 4 (Hierarchical generalized gamma mixture hazard model). If the Dykstra and Laud kernel in Example 3 is combined with a hierarchical vector of generalized gamma CRMs on $\mathbb{Y} = \mathbb{R}^+$, we obtain a hierarchical generalized gamma mixture hazard model. The conditions of (i) in Theorem 1 are met also in this case leading to proper survival functions. The pairwise correlation between survival functions, for any $\ell_1 \neq \ell_2$ and $t_1 \leq t_2$ in $\mathbb{R}^+$ is

$$\text{Cov}(\tilde{S}_{\ell_1}(t_1), \tilde{S}_{\ell_2}(t_2))$$

$$= \exp\left(-\frac{c_0}{\sigma_0} \int_{t_1}^{t_2} [(1 + [(1 + A_2(y))^\sigma - 1]/\sigma)^{\sigma_0} - 1] P_0(dy)\right)$$

(15)
$$\times \left\{\exp\left(-\frac{c_0}{\sigma_0} \int_{0}^{t_1} \left[\left(\sum_{i=1}^{2} [(1 + A_i(y))^\sigma - 1]/\sigma + 1\right)^{\sigma_0} - 1\right] P_0(dy)\right)\right.$$

$$- \exp\left(-\frac{c_0}{\sigma_0}\sum_{i=1}^{2}\int_0^{t_1}\big[(1+[(1+A_i(y))^\sigma-1]/\sigma)^{\sigma_0}-1\big]P_0(\mathrm{d}y)\right)\Bigg\},$$

where $A_i(y) = \alpha(t_i - y)\mathbb{1}_{(0,t_i)}(y)$. This is again a consequence of (ii) in Theorem 1 and calculations detailed in the Supplementary Material.

4. Posterior consistency. In this section, we investigate frequentist posterior consistency of the proposed class of priors. Most of the existing literature on the topic deals with sequences of exchangeable data and other specific forms of dependence (see [21] for an account). But very few results are available for partially exchangeable observations (see, e.g., [1]) and general theorems to study posterior consistency in a multiple samples framework, as the one we are studying here, are still missing. Here, we provide a strategy to face consistency in a partially exchangeable setting and, then, adapt it to show consistency of hierarchical generalized gamma CRMs. Consider the partially exchangeable framework (9) for $\mathbb{R}^+$-valued random elements $X_{\ell,i}$'s and note that the hierarchically dependent mixture hazard model in Definition 1 implies that their distribution is absolutely continuous with respect to the Lebesgue measure. If $(\tilde{f}_1, \ldots, \tilde{f}_d)$ is the vector of random dependent densities associated with $(\tilde{p}_1, \ldots, \tilde{p}_d)$, one has

$$(16) \qquad \tilde{f}_\ell(t) = \tilde{h}_\ell(t)\tilde{S}_\ell(t) = \tilde{h}_\ell(t)\mathrm{e}^{-\tilde{H}_\ell(t)},$$

where $t \mapsto \tilde{H}_\ell(t) = \int_0^t \tilde{h}_\ell(s)\,\mathrm{d}s$ is the cumulative hazard of the ℓth sample, with $\ell = 1, \ldots, d$. Let us denote by $\mathsf{F}_\mathbb{R}$ the space of all probability density functions, with respect to the Lebesgue measure on $\mathbb{R}$. We shall now assume the data from each sequence $(X_{\ell,i})_{i\geq1}$, with $\ell = 1, \ldots, d$, are independently generated from a true and fixed density $f_\ell^{(0)}$ and we, then check whether the posterior distribution of the vector $(\tilde{f}_1, \ldots, \tilde{f}_d)$ accumulates in a neighborhood of $\boldsymbol{f}^{(0)} := (f_1^{(0)}, \ldots, f_d^{(0)})$ in a suitable topology on the product space $\mathsf{F}_\mathbb{R}^d$. In the sequel, we deal with *weak* consistency, hence we endow $\mathsf{F}_\mathbb{R}$ with the *weak* topology and for any $f \in \mathsf{F}_\mathbb{R}$ we denote by $A(f)$ a weak neighborhood of f. Moreover, the space $\mathsf{F}_\mathbb{R}^d$ is naturally endowed with the product topology.

In order to state the main results, we need to introduce some additional notation. First, we let $P_\ell^{(0)}$ denote the probability distribution associated with $f_\ell^{(0)}$ and $P_\ell^{(0),\infty}$ be the infinite product measure. Similarly, $P^{(0),\infty} := P_1^{(0),\infty} \times \cdots \times P_d^{(0),\infty}$. If Π is the prior distribution on $\mathsf{F}_\mathbb{R}^d$ induced by Q, we let $\Pi_{\boldsymbol{n}}(\cdot|X_1, \ldots, X_d)$ be the corresponding posterior and $\boldsymbol{n} = (n_1, \ldots, n_d)$. Hence, the goal is to identify sufficient conditions on the intensities ν_ℓ of the underlying CRMs ($\ell = 0, 1, \ldots, d$) such that, as $n_1, \ldots, n_d \to +\infty$, one has

$$(17) \qquad \Pi_{\boldsymbol{n}}\big(A(\boldsymbol{f}^{(0)})|X_1, \ldots, X_d\big) \to 1, \quad P^{(0),\infty}\text{-a.s.}$$

for any neighborhood $A(\boldsymbol{f}_0)$ of $\boldsymbol{f}^{(0)} = (f_1^{(0)}, \ldots, f_d^{(0)}) \in \mathsf{F}_\mathbb{R}^d$ in the weak topology. If (17) holds true, we say that Π is consistent at $\boldsymbol{f}^{(0)}$. One can then show the following.

THEOREM 2. *Let $(f_1^{(0)}, \ldots, f_d^{(0)})$ be an element of $\mathsf{F}_\mathbb{R}^d$, if for any weak neighborhood $A_\ell(f_\ell^{(0)})$ of $f_\ell^{(0)}$*

$$(18) \qquad \Pi_{\ell,n_\ell}\big(A_\ell(f_\ell^{(0)})|X_\ell\big) := \mathbb{P}\big[\tilde{f}_\ell \in A_\ell(f_\ell^{(0)})|X_\ell\big] \to 1 \quad P_\ell^{(0),\infty}\text{-a.s.}$$

as $n_\ell \to \infty$ for any $\ell = 1, \ldots, d$, then (17) holds true and Π is weakly consistent at $(f_1^{(0)}, \ldots, f_d^{(0)})$.

It is worth remarking two important points related to this result. First, from Theorem 2 it is apparent that showing (17) boils down to proving consistency for each $\tilde{f}_\ell$ given the ℓth sample X_ℓ, and hence, allows us to address the problem in an exchangeable setting. Second, Theorem 2 may be rephrased for vectors of random probability measures $(\tilde{p}_1, \ldots, \tilde{p}_d)$ and its validity is not limited just to random dependent densities.

In view of Theorem 2, in order to show consistency with partially exchangeable arrays, we need to identify conditions for which (18) holds true, for any $\ell = 1, \ldots, d$. Henceforth, we assume that the support of ρ_ℓ is the whole positive real line $\mathbb{R}^+$ and that

$$\int_{\mathbb{R}^+} \rho_\ell(s)\, ds = +\infty$$

for each $\ell = 0, 1, \ldots, d$. Moreover, P_0 is such that its weak support is the whole space $\mathbb{Y}$. We tackle the problem by relying on [9], Theorem 2, which is proved for mixture hazards, and on a result on the support of hierarchical CRMs discussed in Section S2.3 of the Supplementary Material. To fix the notation, $h_\ell^{(0)}$ is the true hazard rate for the ℓ-sample, namely $f_\ell^{(0)}(t) = h_\ell^{(0)}(t) \exp\{-\int_0^t h_\ell^{(0)}(s)\, ds\}$. Let us now recall the following.

THEOREM 3 ([9]). *Let $\tilde{f}_\ell$ be a random density function induced by a random hazard mixture model and denote its distribution by Π_ℓ. If the following conditions hold*:

(i) $f_\ell^{(0)} > 0$ *on* $(0, \infty)$ *and* $\int_{\mathbb{R}^+} \max\{\mathbb{E}\tilde{H}_\ell(t), t\} f_\ell^{(0)}(t)\, dt < +\infty$,
(ii) *either there exists* $r > 0$ *such that* $\liminf_{t \to 0} \tilde{h}_\ell(t)/t^r = \infty$, *a.s., or* $h_\ell^{(0)}(0) > 0$,

a sufficient condition for weak consistency of Π_ℓ at $f_\ell^{(0)}$ is

$$(19) \qquad \Pi_\ell\left(\left\{h : \sup_{t \leq T} |h(t) - h_\ell^{(0)}(t)| < \delta\right\}\right) > 0 \quad \forall T, \delta \in (0, \infty).$$

Theorem 3 provides us with a sufficient condition to prove weak consistency of random densities $\tilde{f}_\ell$. In order to verify condition (ii), one should study the small time behavior of $\tilde{h}_\ell$ otherwise one has to assume that the true hazard rate is such that $h_\ell^{(0)}(0) > 0$ as in [14].

We now focus on the case $\mathbb{Y} = \mathbb{R}^+$, and identify easier sufficient conditions ensuring the validity of Theorem 3(ii); see, for example, Proposition 3 in [9]. We shall do this with hierarchies of the generalized gamma random measure

$$(20) \qquad \rho_0(s) = \frac{1}{\Gamma(1 - \sigma_0)} \frac{e^{-s}}{s^{1+\sigma_0}} \quad \text{and} \quad \rho_\ell(s) \equiv \rho(s) = \frac{1}{\Gamma(1 - \sigma)} \frac{e^{-s}}{s^{1+\sigma}}$$

as $\ell = 1, \ldots, d$, being $\sigma, \sigma_0 \in (0, 1)$. This prior specification will be adopted in the numerical illustrations of Section 7.

PROPOSITION 1. *Let $k(\cdot; \cdot)$ be the Dykstra and Laud kernel and let $\tilde{\mu}_\ell$ be the hierarchical generalized gamma process as in* (20), *with* $\sigma, \sigma_0 \in (0, 1)$ *and P_0 such that $P_0((0, t))/t^{r'} \to C \in (0, \infty)$ as $t \to 0$ for some $r' > 0$. Then the corresponding mixture hazard $\tilde{h}_\ell$ satisfies condition* (ii) *of Theorem 3.*

We can state a theorem concerning marginal consistency of a hierarchical mixture hazard based on generalized gamma processes.

THEOREM 4. *Let $\tilde{h}_\ell$ be a hierarchical mixture hazard as in Proposition 1. Then Π_ℓ is consistent at any $f_\ell^{(0)} \in \mathsf{F}_1$, where F_1 is the set of densities satisfying $\int_{\mathbb{R}^+} \mathbb{E}[\tilde{H}(t)] f_\ell^{(0)}(t)\, dt < \infty$, $h_\ell^{(0)}(0) = 0$ and $h_\ell^{(0)}(t)$ is strictly positive and nondecreasing for any $t > 0$.*

Combining Theorems 2 and 4, we obtain a consistency theorem for Π. With $\mathsf{F}_1^d = \mathsf{X}_1^d \mathsf{F}_1$, one easily deduces the following.

COROLLARY 1. *Let $k(\cdot; \cdot)$ be the Dykstra and Laud kernel and $(\tilde{\mu}_1, \ldots, \tilde{\mu}_d)$ a vector of hierarchical generalized gamma processes that meets the conditions in Proposition 1. Then Π is consistent at any $(f_1^{(0)}, \ldots, f_d^{(0)}) \in \mathsf{F}_1^d$.*

Along similar lines, one may prove consistency for other kernels k, as the ones considered by De Blasi et al. [9].

REMARK 3. If $\sigma = \sigma_0 = 0$, we obtain the gamma process, but condition (ii) is not satisfied in such a case and it should be replaced with a condition on the true hazard rate $(h_\ell^{(0)}(0) > 0)$. Indeed, Drăghici and Ramamoorthi [14] show the consistency under this assumption and they consider a random hazard rate of the following type:

$$\tilde{h}_\ell(t) = \tilde{h}_\ell(0) + \tilde{\mu}_\ell(0, t),$$

where $\tilde{h}_\ell(0)$ is a random element with support on $\mathbb{R}^+$, to consistently estimate the value of the hazard rate in $t = 0$.

5. Random partitions and Bayesian inference. The primary goal we wish to pursue here is the determination of a conditional probability distribution of the vector $(\tilde{\mu}_1, \ldots, \tilde{\mu}_d)$ in (5), given data from a partially exchangeable array and given a suitable collection of latent variables. Indeed, it will be assumed that the data are from an array $\{(X_{\ell,j})_{j \geq 1} : \ell = 1, \ldots, d\}$ as in (9) with

$$(21) \qquad \tilde{p}_\ell((t, +\infty)) = \exp\left(- \int_\mathbb{Y} \int_0^t k(s, y) \, ds \, \tilde{\mu}_\ell(dy)\right)$$

for any $\ell = 1, \ldots, d$ and $t > 0$. The hierarchical specification of $\tilde{\boldsymbol{\mu}} = (\tilde{\mu}_1, \ldots, \tilde{\mu}_d)$, combined with the almost sure discreteness of the base measure $\tilde{\mu}_0$, poses some serious analytical challenges that need to be addressed if one wants to determine Bayesian inferences for survival data modeled through the random probability measure in (21). Hence, from a technical standpoint achieving a posterior characterization is much more difficult than analogous results for exchangeable (see [25]) and for alternative partially exchangeable priors (see, e.g., [27]). Here, we successfully tackle this issue and show that the results we get are not only of theoretical interest per se, but they are also fundamental for devising efficient sampling schemes for an approximate evaluation of Bayesian inferences in this framework. The key is the introduction of two collections of latent variables that describe a sampling procedure somehow reminiscent of the Chinese restaurant franchise scheme introduced in [41] and extended to a more general framework in [4].

5.1. *Latent variables and their partition structure.* For simplicity in the sequel, we will assume that all data are exact. The possible presence of censored observations, with censoring times being independent from all other random components of the model, can be easily accommodated for and this will be explicitly seen in Section 7 through an illustrative example. In order to keep the notation concise, we set $\tilde{\boldsymbol{\mu}} = (\tilde{\mu}_\ell)_\ell$ and $\boldsymbol{X} = (X_\ell)_\ell$. From [24], it is seen that the likelihood function associated to a multiplicative intensity model as the one we are considering here is

$$(22) \qquad \mathscr{L}(\tilde{\boldsymbol{\mu}}; \boldsymbol{X}) = e^{-\sum_{\ell=1}^d \int_\mathbb{Y} K_\ell(y) \tilde{\mu}_\ell(dy)} \prod_{\ell=1}^d \prod_{i=1}^{n_\ell} \int_\mathbb{Y} k(X_{\ell,i}; y) \tilde{\mu}_\ell(dy),$$

where

$$(23) \qquad K_\ell(y) := \sum_{i=1}^{n_\ell} K_{X_{\ell,i}}^{(\ell)}(y) = \sum_{i=1}^{n_\ell} \int_0^{X_{\ell,i}} k(s, y) \, ds$$

for $\ell = 1, \ldots, d$. A considerable simplification of (22) occurs if one removes the integrals by introducing a suitable sequence of latent variables $Y = (Y_\ell)_\ell$. This leads to

$$(24) \qquad \mathscr{L}(\tilde{\mu}; X, Y) = e^{-\sum_{\ell=1}^d \int_{\mathbb{Y}} K_\ell(y) \tilde{\mu}_\ell(dy)} \prod_{\ell=1}^d \prod_{i=1}^{n_\ell} k(X_{\ell,i}; Y_{\ell,i}) \tilde{\mu}_\ell(dY_{\ell,i}).$$

It is worth noting that $Y = (Y_\ell)_\ell$ forms an array of partially exchangeable random elements and (24) suggests they are generated by a discrete random probability measure. Hence, with positive probability there will be tied values in Y both within each sample Y_ℓ and across samples Y_{ℓ_1} and Y_{ℓ_2}, for $\ell_1 \neq \ell_2$. In other terms, Y generates a random partition of the integers $[N] = \{1, \ldots, N\}$, where $N = n_1 + \cdots + n_d$. This can be described by ordering the $Y_{\ell,i}$'s in such a way that $Y_{\ell,i} = Y_{\ell',j}$ entails $\bar{n}_{\ell-1} + i$ and $\bar{n}_{\ell'-1} + j$ are in the same partition set, where $\bar{n}_\ell = n_1 + \cdots + n_\ell$ for each $\ell = 1, \ldots, d$ with the proviso $\bar{n}_0 = 0$. According to this notation, for any partition $\mathcal{C} = \{\bar{C}_1, \ldots, \bar{C}_k\}$ of $[N]$, one can further decompose $\bar{C}_j = \bar{C}_{1,j} \cup \cdots \cup \bar{C}_{d,j}$ with $\bar{C}_{\ell,j} = \{\bar{n}_{\ell-1} + i \in \bar{C}_j : i = 1, \ldots, n_\ell\}$. In the sequel, we denote by Ψ_N the random partition of $[N]$ generated by Y, and by $Y^* := (Y_1^*, \ldots, Y_k^*)$ the k distinct values associated with such a partition. With this notation, $\bar{C}_{\ell,j}$ refers to all elements of the ℓth sample that coincide with Y_j^*. It is easy to realize that the distribution of (X, Y^*, Ψ_N) is equivalent to the distribution of the vector (X, Y), which will be characterized in Theorem 5 below. In order to simplify the notation, we also set

$$Q(X, Y^*) = \prod_{\ell=1}^d \prod_{j=1}^k \prod_{i \in \bar{C}_{\ell,j}} k(X_{\ell,i}; Y_{\ell,i}), \qquad \tau_q^{(\ell)}(u) = \int_0^\infty s^q e^{-su} \rho_\ell(s) \, ds$$

for any $u > 0$, $q \in \mathbb{N}$ and $\ell = 0, 1, \ldots, d$. The subsequent expressions (25) and (27) we determine depend on sums over q and i, and they may appear difficult to evaluate and interpret at a first glance. Nonetheless, a considerable simplification, and additional intuition on these results, is achieved if one resorts to a version of the Chinese restaurant franchise metaphor suited to this particular setting. The franchise is made of d restaurants and the $Y_{\ell,i}$ variables identify the specific dish chosen from the menu by customer i at the ℓth restaurant. The values $Y_1^*, \ldots, Y_k^*$ are the k distinct dishes selected by the N customers at the d restaurants of the franchise and $n_{\ell,j} = \text{card}\{i : Y_{\ell,i} = Y_j^*\}$ is the number of customers being served dish j at the ℓth restaurant. Of course, one may have that $\{i : Y_{\ell,i} = Y_j^*\} = \varnothing$, in which case $n_{\ell,j} = 0$. With reference to previous notation, the set $C_{\ell,j} := \{i : Y_{\ell,i} = Y_j^*\}$ plays the role of $\bar{C}_{\ell,j}$, whose elements are equal to those of $C_{\ell,j}$ after subtracting the additive constant $\bar{n}_{\ell-1}$. At this stage it is then convenient to introduce latent variables $T_\ell := (T_{\ell,1}, \ldots, T_{\ell,n_\ell})$ where, for each $\ell = 1, \ldots, d$, $T_{\ell,j}$ is the label of the table where the jth customer of restaurant ℓ is sitting and with the convention that customers seating at the same tables are being served the same dish. The $T_{\ell,i}$'s are generated by a discrete random probability measure, and hence admit ties, namely $\{T_{\ell,i} : i \in C_{\ell,j}\}$ display $i_{\ell,j} \leq n_{\ell,j}$ distinct values $T_{\ell,j,1}^*, \ldots, T_{\ell,j,i_{\ell,j}}^*$ that induce a partition of $[n_{\ell,j}] = \{1, \ldots, n_{\ell,j}\}$ into $i_{\ell,j}$ sets

$$C_{\ell,j,t} = \{i \in C_{\ell,j} : T_{\ell,i} = T_{\ell,j,t}^*\} \quad (t = 1, \ldots, i_{\ell,j})$$

with $q_{\ell,j,t} = \text{card}(C_{\ell,j,t})$ such that $\sum_{t=1}^{i_{\ell,j}} q_{\ell,j,t} = n_{\ell,j}$. Note that $i_{\ell,j} = 0$ when $n_{\ell,j} = 0$ and we, thus, agree that $\sum_{t=1}^0 a_t = 0$ for any sequence $(a_t)_{t\geq 1}$. We are now ready to prove the main results of this section that can be read in terms of the metaphor just outlined.

THEOREM 5. *The probability distribution of (X, Y^*) and of Ψ_N equals*

$$\pi(X, Y^*, \Psi_N)$$

$$\propto Q(X, Y^*) c_0^k e^{-c_0 \int_{\mathbb{Y}} \psi^{(0)}(\sum_{\ell=1}^d \psi^{(\ell)}(K_\ell(y))) P_0(\mathrm{d}y)}$$

(25)
$$\times \sum_i \sum_q \prod_{j=1}^k \tau_{i_{\bullet j}}^{(0)} \left(\sum_{\ell=1}^d \psi^{(\ell)}(K_\ell(Y_j^*)) \right) P_0(\mathrm{d}Y_j^*)$$

$$\times \prod_{\ell=1}^d \prod_{j=1}^k \frac{1}{i_{\ell,j}!} \binom{n_{\ell,j}}{q_{\ell,j,1}, \ldots, q_{\ell,j,i_{\ell,j}}} \prod_{t=1}^{i_{\ell,j}} \tau_{q_{\ell,j,t}}^{(\ell)}(K_\ell(Y_j^*)),$$

where the sum with respect to q runs over all the vectors of positive integers $\{q_{\ell,j,t}\}$ such that $\sum_{t=1}^{i_{\ell,j}} q_{\ell,j,t} = n_{\ell,j}$ for any $\ell = 1, \ldots, d$ and $j = 1, \ldots, k$ and $i = \{i_{\ell,j}\}$ is through the set of all integers $i_{\ell,j} \in \{1, \ldots, n_{\ell,j}\}$, with $i_{\bullet j} = \sum_{\ell=1}^d i_{\ell,j}$, for any $j = 1, \ldots, k$.

This result is the backbone of theoretical and computational developments in the remainder of the paper. For example, one can deduce from (25) the following augmented joint distribution:

$$\pi(X, Y^*, T, \Psi_N)$$

$$\propto c_0^k Q(X, Y^*) e^{-c_0 \int_{\mathbb{Y}} \psi^{(0)}(\sum_{\ell=1}^d \psi^{(\ell)}(K_\ell(y))) P_0(\mathrm{d}y)}$$

(26)
$$\times \prod_{j=1}^k \tau_{i_{\bullet j}}^{(0)} \left(\sum_{\ell=1}^d \psi^{(\ell)}(K_\ell(Y_j^*)) \right) P_0(\mathrm{d}Y_j^*)$$

$$\times \prod_{\ell=1}^d \prod_{j=1}^k \prod_{t=1}^{i_{\ell,j}} \tau_{q_{\ell,j,t}}^{(\ell)}(K_\ell(Y_j^*)) P_0(\mathrm{d}T_{\ell,j,t}^*),$$

where $i = \{i_{\ell,j}\}$ and $q = \{q_{\ell,j,t}\}$ are now fixed. Note that, since the specific table's label is not relevant, for simplicity and with no loss of generality we have assumed that each table's label still takes values in $\mathbb{Y}$. This formula provides some nice intuition on the Chinese restaurant franchise metaphor briefly discussed before Theorem 5. Indeed, if we agree that $D_{-(\ell,i)}$ is the vector (X, Y^*, T, q, i) after the removal of $(X_{\ell,i}, Y_{\ell,i}, T_{\ell,i})$, one has that the dish chosen by the ith customer in the ℓth restaurant and the table where she seats are determined through the following distribution

$$\mathbb{P}[Y_{\ell,i} = Y_j^*, T_{\ell,i} = T_{\ell,j,t}^* | D_{-(\ell,i)}] \propto k(X_{\ell,i}; Y_j^*) \frac{\tau_{q_{\ell,j,t}+1}^{(\ell)}(K_\ell(Y_j^*))}{\tau_{q_{\ell,j,t}}^{(\ell)}(K_\ell(Y_j^*))},$$

which corresponds to the customer choosing the jth dish already ordered from the menu and seating at the already existing tth table, whereas

$$\mathbb{P}[Y_{\ell,i} = Y_j^*, T_{\ell,i} \notin \mathcal{T}_{-(\ell,i)} | D_{-(\ell,i)}] \propto k(X_{\ell,i}; Y_j^*) \frac{\tau_{i_{\bullet,j}+1}^{(0)}(\sum_{r=1}^d \psi^{(r)}(Y_j^*))}{\tau_{i_{\bullet,j}}^{(0)}(\sum_{r=1}^d \psi^{(r)}(Y_j^*))}$$

is the probability that the customer still chooses the jth dish though decides to sit at a new table whose label will be drawn from P_0. Finally, the customer may choose a new dish, and then sit at a new table with probability

$$\mathbb{P}[Y_{\ell,i} \notin \mathcal{Y}_{-(\ell,i)}, T_{\ell,i} \notin \mathcal{T}_{-(\ell,i)} | D_{-(\ell,i)}]$$

$$\propto c_0 \int_{\mathbb{Y}} k(X_{\ell,i}; y) \tau_1^{(\ell)}(K_\ell(y)) \tau_1^{(0)} \left(\sum_{r=1}^{d} \psi^{(r)}(K_r(y)) \right) P_0(\mathrm{d}y)$$

and the label of both the new dish and the new table are generated from P_0. This structure of the Chinese restaurant franchise will be exploited in the algorithm that will be used for determining Bayesian inferences. For details, see the Supplementary Material.

Theorem 5 is also relevant for deducing the distribution of the random partition Ψ_N conditionally on the other variables of the model.

COROLLARY 2. *Let $\mathcal{C} = \{\bar{C}_1, \ldots, \bar{C}_k\}$ denote a partition of $[N]$ into k sets such that $\bar{C}_j = \bar{C}_{1,j} \cup \cdots \cup \bar{C}_{d,j}$. Moreover, we let $\mathrm{card}(\bar{C}_j) = n_j \geq 1$ and $\mathrm{card}(\bar{C}_{\ell,j}) = n_{\ell,j} \geq 0$. If Ψ_N denotes the random partition of $[N]$ induced by $\mathbf{Y}$, then $\mathbb{P}[\Psi_N = \mathcal{C} | \mathbf{X}]$ is proportional to*

$$
c_0^k \sum_{\mathbf{i}} \sum_{\mathbf{q}} \left\{ \prod_{\ell=1}^{d} \prod_{j=1}^{k} \frac{1}{i_{\ell,j}!} \binom{n_{\ell,j}}{q_{\ell,j,1}, \ldots, q_{\ell,j,i_{\ell,j}}} \right\}
$$

(27)
$$
\times \prod_{j=1}^{k} \int_{\mathbb{Y}} \left\{ \prod_{\ell=1}^{d} \prod_{i \in \bar{C}_{\ell,j}} k(X_{\ell,i-\bar{n}_{\ell-1}}; y) \right\} \tau_{i \bullet j}^{(0)} \left(\sum_{\ell=1}^{d} \psi^{(\ell)}(K_\ell(y)) \right)
$$

$$
\times \prod_{\ell=1}^{d} \prod_{t=1}^{i_{\ell,j}} \tau_{q_{\ell,j,t}}^{(\ell)}(K_\ell(y)) P_0(\mathrm{d}y),
$$

where the sums with respect to $\mathbf{q}$ and $\mathbf{i}$ are as in Theorem 5.

At this point, one may wonder whether there exist CRMs such that

(28)
$$
\pi(\mathbf{Y}^*, \mathbf{T}, \Psi_N | \mathbf{X}) = \int \pi(\mathbf{Y}^*, \mathbf{T}, \Psi_N | \mathbf{X}, \boldsymbol{\mu}) \mathbb{P}(\mathrm{d}\boldsymbol{\mu} | \mathbf{X}),
$$

the answer is provided by the following result, which is based on Corollary 2.

THEOREM 6. *Let $m_0 \sim \mathrm{CRM}(\nu_0)$ and $m_{\ell,y} \sim \mathrm{CRM}(\nu_{\ell,y})$ be independent and such that*

$$
\nu_0(\mathrm{d}s, \mathrm{d}y) = c_0 e^{-\sum_{\ell=1}^{d} \psi^{(\ell)}(K_\ell(y))} \rho_0(s) \, \mathrm{d}s \, P_0(\mathrm{d}y),
$$

$$
\nu_{\ell,y}(\mathrm{d}s, \mathrm{d}w) = e^{-K_\ell(y)w} \rho_\ell(s) \, \mathrm{d}s \, P_0(\mathrm{d}w).
$$

Then

$$
\pi(\mathbf{Y}^*, \mathbf{T}, \Psi_N | \mathbf{X}, m_0, \boldsymbol{m})
$$

(29)
$$
\propto \prod_{\ell=1}^{d} \prod_{j=1}^{k} \left\{ \prod_{i \in C_{\ell,j}} k(X_{\ell,i}, Y_j^*) \right\} \prod_{t=1}^{i_{\ell,j}} m_0(\mathrm{d}Y_j^*) \prod_{r \in C_{\ell,j,t}} m_{\ell,Y_j^*}(\mathrm{d}T_{\ell,j,t}^*),
$$

where $\boldsymbol{m}$ denotes the vector containing the m_{ℓ,Y_j^*}'s.

We have previously defined the random partition Ψ_N in such a way that there is a one to one correspondence between the two vectors $(\mathbf{Y}^*, \Psi_N)$ and $\mathbf{Y}$. Henceforth, we equivalently write $\mathbf{Y}$ instead of $(\mathbf{Y}^*, \Psi_N)$.

5.2. *A posterior characterization.* The result in Theorem 5, combined with the related augmented probability distribution in (26), paves the way to the determination of the posterior distribution of $\tilde{\boldsymbol{\mu}} = (\tilde{\mu}_1, \ldots, \tilde{\mu}_d)$, given the data and the latent variables. This is described in the next theorem, which shows that a structural conjugacy property holds true and the vector $\tilde{\boldsymbol{\mu}}$ is still hierarchical *a posteriori*. In order to state the result, let $I_1, \ldots, I_k$ be independent nonnegative random variables with I_j having density function

$$(30) \qquad f_j(s) \propto s^{i \cdot j} e^{-s \sum_{\ell=1}^{d} \psi^{(\ell)}(K_\ell(Y_j^*))} \rho_0(s) \, ds$$

and $\eta_0^* \sim \mathrm{CRM}(\nu_0^*)$ where

$$(31) \qquad \nu_0^*(ds, dy) = e^{-s \sum_{\ell=1}^{d} \psi^{(\ell)}(K_\ell(y))} \rho_0(s) \, ds c_0 P_0(dy).$$

Finally, set $\tilde{\mu}_0^* = \eta_0^* + \sum_{j=1}^{k} I_j \delta_{Y_j^*}$ and correspondingly

$$(32) \qquad \tilde{\nu}_\ell^*(ds, dy) = e^{-s K_\ell(y)} \rho_\ell(s) \, ds \, \tilde{\mu}_0^*(dy)$$

for each $\ell = 1, \ldots, d$.

THEOREM 7. *The posterior distribution of $\tilde{\boldsymbol{\mu}}$, conditional on $(\boldsymbol{X}, \boldsymbol{Y}, \boldsymbol{T})$, equals the distribution of the vector of CRMs*

$$(33) \qquad (\tilde{\mu}_1^*, \ldots, \tilde{\mu}_d^*) + \left(\sum_{j=1}^{k_1} \sum_{t=1}^{i_{1,j}} J_{1,j,t} \delta_{Y_{1,j}^*}, \ldots, \sum_{j=1}^{k_d} \sum_{t=1}^{i_{d,j}} J_{d,j,t} \delta_{Y_{d,j}^*} \right),$$

where $\tilde{\mu}_\ell^* | \tilde{\mu}_0^* \overset{\mathrm{ind}}{\sim} \mathrm{CRM}(\tilde{\nu}_\ell^*)$, for $\ell = 1, \ldots, d$, and the jumps $J_{\ell,j,t}$ are independent and nonnegative random elements with corresponding density functions

$$(34) \qquad f_{\ell,j,t}(s) \propto s^{q_{\ell,j,t}} e^{-s K_\ell(Y_{\ell,j}^*)} \rho_\ell(s) \, ds.$$

Moreover, the random elements $\tilde{\boldsymbol{\mu}}^* = (\tilde{\mu}_1^*, \ldots, \tilde{\mu}_d^*)$, η_0^*, $(I_1, \ldots, I_k)$ and $\{J_{\ell,j,t}\}$ are mutually independent.

It is worth stressing that according to Theorem 7, the random measures $\tilde{\mu}_\ell^* + \sum_{j=1}^{k_\ell} \sum_{t=1}^{i_{\ell,j}} J_{\ell,j,t} \delta_{Y_{\ell,j}^*}$ are conditionally independent CRMs, given $\tilde{\mu}_0^*$. The main differences with respect to the prior (5) are the presence of the jumps $J_{\ell,j,t}$ at fixed locations identified by the distinct values $Y_{\ell,j}^*$ of the latent variables $\boldsymbol{Y}_\ell$ and the exponential updating of the conditional Lévy intensity as described in (32). Moreover, unlike (5) the CRM $\tilde{\mu}_0^*$ at the top hierarchy does also have jumps at fixed locations corresponding to the overall distinct latent variables' values $Y_1^*, \ldots, Y_k^*$. Hence the hierarchical structure is preserved also *a posteriori* and a property of structural conjugacy holds true. Note that the posterior characterization given in Theorem 7 shares some interesting features with the one of [4], Theorem 10. Indeed, at the CRM level, the posteriors exhibit structural analogies displaying an updated CRM independent of positive fixed jumps I_j's, that are shared across the groups, and of positive fixed jumps $J_{\ell,j,t}$'s, that are specific to group ℓ. Otherwise, the posteriors are very different in distribution, given the different transformation of the CRMs they rely on. This somehow surprising and interesting finding is reminiscent of the structural analogies pointed out by [29] for the simple exchangeable case.

A posterior characterization, as the one in Theorem 7, is the starting point for evaluating Bayesian inferences of interest on functionals of $(\tilde{p}_1, \ldots, \tilde{p}_d)$. For example, one can resort to (33) for determining a posterior estimate of the sample-specific survival functions, in the following.

COROLLARY 3. *For any $t > 0$, the posterior estimate of the survival function $\tilde{S}_\ell(t)$ with respect to a quadratic loss is*

$$
\mathbb{E}\big[\tilde{S}_\ell(t)|\boldsymbol{X}, \boldsymbol{Y}, \boldsymbol{T}\big]
$$

$$
= \exp\Big(-c_0 \int_{\mathbb{Y}} \psi_*^{(0)}\big(\psi_*^{(\ell)}(K_t^{(\ell)}(y))\big) P_0(\mathrm{d}y)\Big)
$$

(35)

$$
\times \prod_{j=1}^{k} \frac{\int_0^\infty e^{-s\{\psi_*^{(\ell)}(K_t^{(\ell)}(Y_j^*))+\sum_{h=1}^{d}\psi^{(h)}(K_h(Y_j^*))\}} s^{i_{\bullet j}} \rho_0(s)\,\mathrm{d}s}{\int_0^\infty e^{-s\sum_{h=1}^{d}\psi^{(h)}(K_h(Y_j^*))} s^{i_{\bullet j}} \rho_0(s)}
$$

$$
\times \prod_{j=1}^{k_\ell} \prod_{t=1}^{i_{\ell,j}} \frac{\int_0^\infty e^{-s(K_t^{(\ell)}(Y_{\ell,j}^*)+K_\ell(Y_{\ell,j}^*))} s^{q_{\ell,j,t}} \rho_\ell(s)\,\mathrm{d}s}{\int_0^\infty e^{-sK_\ell(Y_{\ell,j}^*)} s^{q_{\ell,j,t}} \rho_\ell(s)\,\mathrm{d}s},
$$

where

$$
\psi_*^{(0)}(u) = \int_0^\infty (1 - e^{-su}) e^{-s\sum_{\ell=1}^{d}\psi^{(\ell)}(K_\ell(y))} \rho_0(s)\,\mathrm{d}s,
$$

$$
\psi_*^{(\ell)}(u) = \int_0^\infty (1 - e^{-su}) e^{-sK_\ell(y)} \rho_\ell(s)\,\mathrm{d}s, \quad \ell = 1, \dots, d.
$$

The posterior expectation of $\tilde{S}_\ell$, as described in Corollary 3, is conditional on the data $\boldsymbol{X}$ and on all the latent variables. Therefore, in order to obtain a posterior estimate of the survival function, one marginalizes (35) with respect to the posterior distribution of $(\boldsymbol{Y}, \boldsymbol{T})$, given $\boldsymbol{X}$. This goal may be achieved resorting to a Gibbs sampler, since such a marginalization cannot be handled analytically, and this will be the subject of Section 6.1. Analogously, one can obtain the posterior expectation of the hazard rate for each group of survival times.

COROLLARY 4. *For any $t > 0$, the posterior estimate of the hazard rate $\tilde{h}_\ell(t)$ conditionally given $\boldsymbol{X}, \boldsymbol{Y}, \boldsymbol{T}$, under a square loss function is*

$$
\int_{\mathbb{Y}} k(t; y) \int_0^\infty s e^{-sK_\ell(y)} \rho_\ell(s)\,\mathrm{d}s \int_0^\infty w e^{-w\sum_{h=1}^{d}K_h(y)} \rho_0(w)\,\mathrm{d}w c_0 P_0(\mathrm{d}y)
$$

$$
+ \sum_{j=1}^{k} k(t; Y_j^*) \int_0^\infty s e^{-sK_\ell(Y_j^*)} \rho_\ell(s)\,\mathrm{d}s \int_0^\infty w f_j(w)\,\mathrm{d}w
$$

$$
+ \sum_{j=1}^{k_\ell} \sum_{t=1}^{i_{\ell,j}} k(t; Y_{\ell,j}^*) \int_0^\infty w f_{\ell,j,t}(w)\,\mathrm{d}w.
$$

6. Marginal and conditional samplers. We now rely on the theoretical finding of the previous sections to describe a marginal and a conditional sampler, which can be used, for instance, to obtain posterior estimates of the survival functions. More precisely in Section 6.1, we develop a MCMC procedure to estimate dependent survival functions, based on Theorem 7 and Corollary 3. We in fact use the output of such an MCMC procedure to estimate the posterior expected values of the random survival functions, marginalizing out the CRMs. Hence in Section 6.1 we devise a so-called *marginal* algorithm. On the other hand, we also describe a *conditional* strategy in Section 6.2, which allows us to simulate the trajectories of the hierarchical CRMs.

6.1. *A marginal MCMC sampler.* Though here we will focus on a special case where each $\tilde{\mu}_\ell$ is a generalized gamma process, more specifically $\rho_\ell(s) = e^{-s} s^{-1-\sigma}/\Gamma(1-\sigma)$ for $\ell = 1, \ldots, d$ and $\rho_0(s) = e^{-s} s^{-1-\sigma_0}/\Gamma(1-\sigma_0)$, our analysis can be extended to any other choice of Lévy intensities that satisfy the conditions (i)–(ii) in Theorem 1. We further assume that $\mathbb{Y} = \mathbb{R}^+$ and set $k(t; y) := \mathbb{1}_{(0,t]}(y)\alpha(y)$ for some positive right-continuous real-valued function $\alpha(\cdot)$. Note that when $\sigma = \sigma_0 = 0$, each random hazard rate is an extended gamma process, given $\tilde{\mu}_0$, with parameters $(\tilde{\mu}_0, \alpha)$. The extended gamma process was introduced in [15] to model monotone increasing hazard rates and it has been extensively used in an exchangeable setting. We will further assume that the function $\alpha(y) \equiv \alpha$ is constant and we specify a prior for α. We are able to deal with the general case $d \geq 2$, without analytical difficulties with respect the situation in which $d = 2$. One can now specialize Corollary 3 to this case and get the following result.

COROLLARY 5. *Let $\rho_\ell(s) = e^{-s} s^{-1-\sigma}/\Gamma(1-\sigma)$ for $\ell = 1, \ldots, d$, $\rho_0(s) = e^{-s} s^{-1-\sigma_0}/ \Gamma(1-\sigma_0)$, and $k(t; y) := \alpha \mathbb{1}_{(0,t]}(y)$. Then the posterior expected value $\mathbb{E}[\tilde{S}_\ell(t)|\boldsymbol{X}, \boldsymbol{Y}, \boldsymbol{T}]$ equals*

$$\exp\left(-\frac{c_0}{\sigma_0} \int_{\mathbb{Y}} F_\ell(y) P_0(\mathrm{d}y)\right)$$

$$\times \prod_{j=1}^{k}\left\{1 + \frac{[1 + K_\ell(Y_j^*) + K_t^{(\ell)}(Y_j^*)]^\sigma - [1 + K_\ell(Y_j^*)]^\sigma}{\sum_{h=1}^{d}[(K_h(Y_j^*) + 1)^\sigma - 1]}\right\}^{-(i_{\bullet j} - \sigma_0)}$$

$$\times \prod_{j=1}^{k_\ell}\left\{1 + \frac{K_t^{(\ell)}(Y_{\ell,j}^*)}{1 + K_\ell(Y_{\ell,j}^*)}\right\}^{-(n_{\ell,j} - i_{\ell,j}\sigma)}$$

being

$$F_\ell(y) = \left[\sum_{h=1}^{d} \frac{(K_h(y) + 1)^\sigma - 1}{\sigma} + 1\right]^{\sigma_0}$$

$$\times \left\{\left[1 + \frac{[1 + K_\ell(y) + K_t^{(\ell)}(y)]^\sigma - [1 + K_\ell(y)]^\sigma}{\sigma[\sum_{h=1}^{d} \frac{(K_h(y)+1)^\sigma - 1}{\sigma} + 1]}\right]^{\sigma_0} - 1\right\}$$

for each $\ell = 1, \ldots, d$.

As far as the hazard is concerned, an application of Corollary 4 in the case of hierarchies of generalized gamma process leads to the following.

COROLLARY 6. *Let $\rho_\ell(s) = e^{-s} s^{-1-\sigma}/\Gamma(1-\sigma)$ for $\ell = 1, \ldots, d$, $\rho_0(s) = e^{-s} s^{-1-\sigma_0}/ \Gamma(1-\sigma_0)$, and $k(t; y) := \alpha \mathbb{1}_{(0,t]}(y)$, then the posterior expected value $\mathbb{E}[\tilde{h}_\ell(t)|\boldsymbol{X}, \boldsymbol{Y}, \boldsymbol{T}]$ equals*

$$\int_{\mathbb{Y}} \frac{k(t; y)}{(1 + \sum_{h=1}^{d} K_h(y))^{1-\sigma_0}(1 + K_\ell(y))^{1-\sigma}} c_0 P_0(\mathrm{d}y)$$

(36)
$$+ \sum_{j=1}^{k} \frac{k(t; Y_j^*)(i_{\bullet j} - \sigma_0)}{(1 + \sigma^{-1}\sum_{h=1}^{d}[(1 + K_h(Y_j^*))^\sigma - 1])(1 + K_\ell(Y_j^*))^{1-\sigma}}$$

$$+ \sum_{j=1}^{k_\ell} \sum_{t=1}^{i_{\ell,j}} \frac{k(t; Y_{\ell,j}^*)(q_{\ell,j,t} - \sigma)}{(1 + K_\ell(Y_{\ell,j}^*))}$$

for any $\ell = 1, \ldots, d$.

It is now immediate to check that, with our choices of the kernel $k(t; y)$ and of the Lévy intensities ρ_ℓ, one has

$$K_t^{(\ell)}(y) = \alpha(t - y)\mathbb{1}_{(0,t]}(y),$$

$$K_\ell(y) = \alpha \sum_{\{i:y \le X_{\ell,i}\}} X_{\ell,i} - y\alpha \cdot \#\{i : y \le X_{\ell,i}\}.$$

An approximate evaluation of the posterior estimate $\mathbb{E}[\tilde{S}_\ell(t)|X]$ of the survival function $\tilde{S}_\ell$ may be obtained by means of an MCMC procedure. At the same time, one can obtain a posterior estimate of the hazard rate $\tilde{h}_\ell(t)$. To this end, the full conditional distributions of the latent variables must be identified. The detailed description of the Gibbs sampler and all the full conditional distributions is reported in Section S5. To complete the picture, we assume α and c_0 are independent and further specify the following priors:

(37) $\alpha \sim \mathrm{Ga}(a, b), \qquad c_0 \sim \mathrm{Ga}(a_0, b_0).$

For computational convenience, σ and σ_0 are assumed to be fixed values.

6.2. *Conditional sampler.* The key distinctive feature of the MCMC procedure detailed in previous Section 6.1 is the marginalization with respect to the mixing dependent CRMs $\tilde{\mu}_\ell$'s. This is very useful when one is mainly interested in determining estimates of the survival functions, at any point $t > 0$, as posterior expected values of $\tilde{S}_\ell(t)$'s; see Corollary 5. On the contrary, in the present section we are going to develop a *conditional* algorithm that generates trajectories of the $\tilde{\mu}_\ell$'s from their respective posterior distributions. Our strategy relies on Theorem 7 and a proper adaptation of the algorithm by Wolpert and Ickstadt [42]. A conditional sampler is very useful for many practical reasons, and here we confine ourselves to just mentioning that: (i) it allows to estimate the actual posterior distribution of the survival functions or, equivalently, of the hazard rates at any time point t; (ii) it yields estimates of the posterior credible intervals for the estimated quantities that are more reliable than those arising from marginal samplers.

As for (i), if $\mathbf{\Phi}_1, \ldots, \mathbf{\Phi}_N$ denote N posterior samples of the latent vector $\mathbf{\Phi}$ obtained through an MCMC procedure, and if in addition $\tilde{\mu}_\ell^{(i,1)}, \ldots, \tilde{\mu}_\ell^{(i,M)}$ are M trajectories of $\tilde{\mu}_\ell$, conditionally given $\mathbf{\Phi}_i, X$, then the posterior distribution function of survival function $\tilde{S}_\ell(t)$, at t, is approximated by

(38) $$\mathbb{P}(\tilde{S}_\ell(t) \le s|X) \approx \frac{1}{NM} \sum_{i=1}^{N} \sum_{k=1}^{M} \mathbb{1}_{\{\tilde{S}_\ell^{(i,k)}(t) \le s\}},$$

where

$$\tilde{S}_\ell^{(i,k)}(t) := \exp\left(-\int_0^t \int_{\mathbb{Y}} k(s; y)\tilde{\mu}_\ell^{(i,k)}(\mathrm{d}y)\, \mathrm{d}s\right).$$

See Section S6.1 for details on the derivation of (38). Being provided with the approximation of the posterior distribution of the survival functions (38), it is now clear one can also address the aforementioned point (ii).

Hence, we need to devise an algorithm that samples the trajectories of the $\tilde{\mu}_\ell$'s from their posterior distribution and plug them in (38). Having fixed $\ell = 1, \ldots, d$, the computational procedure is based on Theorem 7 and a nontrivial adaptation of the algorithm developed by Wolpert and Ickstadt [42]. The technique suggested in [42] is easier to implement in the presence of nonhomogeneous CMRs, as those considered here, if compared to an alternative and popular sampler yielded by the Ferguson and Klass [18] representation. Nonetheless, the jumps of the CRMs are not sampled in a decreasing order and this may be a serious drawback

when one wants to obtain an approximate draw of a CRM by discarding the infinite number of jumps whose size does not exceed a fixed threshold, $\varepsilon > 0$ say. We have been able to successfully address this issue for the CRMs

$$(39) \qquad \eta_0^* = \sum_{h \geq 1} J_h^{(0)} \delta_{y_h^{(0)}}, \qquad \tilde{\mu}_\ell^* = \sum_{h \geq 1} J_h^{(\ell)} \delta_{y_h^{(\ell)}},$$

for $\ell = 1, \ldots, d$, where $y_h^{(0)} \overset{\text{i.i.d.}}{\sim} P_0$ and $y_h^{(\ell)} | \tilde{\mu}_0^* \overset{\text{i.i.d.}}{\sim} \tilde{\mu}_0^* / \tilde{\mu}_0^*(\mathbb{Y})$; see Theorem 7. Let now $S_h^{(0)}$ and $S_h^{(\ell)}$ be the points of a unit rate Poisson process, for $h \geq 1$, namely $S_h^{(0)} - S_{h-1}^{(0)}$ and $S_h^{(\ell)} - S_{h-1}^{(\ell)}$ are independent random variables whose distribution is negative exponential with unit mean. Hence, according to [42] the jumps $J_h^{(0)}$ and $J_h^{(\ell)}$ in (39) are such that

$$(40) \qquad S_h^{(0)} = c_0 \int_{J_h^{(0)}}^{\infty} \mathrm{e}^{-s \sum_{\ell=1}^d \psi_\ell(K_\ell(y_h^{(0)}))} \rho_0(s) \, \mathrm{d}s,$$

$$(41) \qquad S_h^{(\ell)} = \tilde{c} \int_{J_h^{(\ell)}}^{\infty} \mathrm{e}^{-s K_\ell(y_h^{(\ell)})} \rho_\ell(s) \, \mathrm{d}s,$$

where $\tilde{c} := \sum_{h \geq 1} J_h^{(0)} \delta_{y_h^{(0)}} + \sum_{j=1}^k I_j \delta_{Y_j^*}$ is the total mass of $\tilde{\mu}_0^*$. One can then show the following.

PROPOSITION 2. *The sequences of nonnegative random variables* $(\tilde{J}_h^{(0)})_{h \geq 1}$ *and* $(\tilde{J}_h^{(\ell)})_{h \geq 1}$, *for* $\ell = 1, \ldots, d$, *defined through the equations*

$$S_h^{(0)} = c_0 \int_{\tilde{J}_h^{(0)}}^{\infty} \rho_0(s) \, \mathrm{d}s, \qquad S_h^{(\ell)} = \tilde{c} \int_{\tilde{J}_h^{(\ell)}}^{\infty} \rho_\ell(s) \, \mathrm{d}s$$

are monotonically decreasing and dominate $(J_h^{(0)})_{h \geq 1}$ *and* $(J_h^{(\ell)})_{h \geq 1}$ *in* (40)–(41), *namely* $J_h^{(0)} \leq \tilde{J}_h^{(0)}$ *and* $J_h^{(\ell)} \leq \tilde{J}_h^{(\ell)}$, *for any* $h \geq 1$ *and* $\ell = 1, \ldots, d$, *almost surely.*

In order to make precise the use of this result, let us focus attention on η_0^* in (39). If one wants to determine a finite-sum approximation of η_0^*, one can fix $\varepsilon > 0$ and discard jumps whose size is smaller than ε by setting the truncation level H^ε as the minimum value of h such that the additional jump satisfies $\tilde{J}_h^{(0)} \leq \varepsilon$, that is,

$$(42) \qquad \eta_0^* \approx \sum_{h=1}^{H^\varepsilon} J_h^{(0)} \delta_{y_h^{(0)}}.$$

By virtue of Proposition 2, such a procedure guarantees that the heights of the discarded jumps $J_h^{(0)}$'s in (42) are smaller than ε. Similar arguments apply to $\tilde{\mu}_\ell^*$.

We now have all the necessary ingredients to describe the conditional algorithm.

(1) Generate $\tilde{\mu}_0$ from its posterior distribution, which is described right before Theorem 7, namely proceed as follows:

(1.a) Generate the random jumps I_j, as $j = 1, \ldots, k$, whose density is proportional to (30);

(1.b) generate η_0^* using the algorithm developed by Wolpert and Ickstadt [42], that is, fix a threshold level $\varepsilon > 0$ and proceed with the following steps:

– generate the atom of the CRM $y_h^{(0)} \sim P_0$;
– generate the waiting times $S_h^{(0)}$ of a standard Poisson process, that is to say $S_h^{(0)} - S_{h-1}^{(0)}$ are independent and identically distributed exponential random variables with unit mean;

– determine the jump $J_h^{(0)}$ by inverting the upper tail of the Lévy intensity (31), that is, by solving (40);

– stop the procedure at $H_0^\varepsilon := \min\{h : \tilde{J}_h^{(0)} \leq \varepsilon\}$ where the auxiliary jumps $\tilde{J}_h^{(0)}$ are determined according to Proposition 2.

(1.c) Evaluate an approximate draw of $\tilde{\mu}_0^*$

$$\tilde{\mu}_{0,\varepsilon}^* = \sum_{h=1}^{H_0^\varepsilon} J_h^{(0)} \delta_{y_h^{(0)}} + \sum_{j=1}^{k} I_j \delta_{Y_j^*}.$$

(2) Generate $\tilde{\mu}_\ell$, given $(\tilde{\mu}_0^*, X)$, as follows.

(2.a) Generate the $J_{\ell,j,t}$'s from the density in (34);

(2.b) with $\varepsilon > 0$ fixed, generate the $\tilde{\mu}_\ell^*$'s using the posterior representation in Theorem 7 and the algorithm by Wolpert and Ickstadt [42], namely:

– generate the atoms of the CRMs $y_h^{(\ell)} \sim P_{0,\varepsilon}^* = \tilde{\mu}_{0,\varepsilon}^* / \tilde{\mu}_{0,\varepsilon}^*(\mathbb{Y})$;

– generate the waiting times $S_h^{(\ell)}$ of a standard Poisson process, that is to say $S_h^{(\ell)} - S_{h-1}^{(\ell)}$ are independent and identically distributed exponential random variables with unit mean;

– determine the jump $J_h^{(\ell)}$ by inverting the upper tail of the Lévy intensity (41);

– stop the procedure at $H_\ell^\varepsilon := \min\{h : \tilde{J}^{(\ell)} \leq \varepsilon\}$, where the auxiliary jumps $\tilde{J}_h^{(\ell)}$ are determined according to Proposition 2;

(2.c) evaluate an approximate draw from the posterior of $\tilde{\mu}_\ell$, by putting

$$\tilde{\mu}_\ell \approx \sum_{h=1}^{H_\ell^\varepsilon} J_h^{(\ell)} \delta_{y_h^{(\ell)}} + \sum_{j=1}^{k_\ell} \sum_{t=1}^{i_{\ell,j}} J_{\ell,j,t} \delta_{Y_{\ell,j}^*}.$$

For the numerical experiments of Section 7, we have implemented the conditional algorithm when $\tilde{\mu}_\ell$ is such that $\rho_\ell(s) = (\Gamma(1-\sigma))^{-1} e^{-s}/s^{1+\sigma}$ for $\ell = 1, \ldots, d$, and $\rho_0(s) = (\Gamma(1-\sigma_0))^{-1} e^{-s}/s^{1+\sigma_0}$, that is, the CRMs are generalized gamma processes exactly as in the previous section. For the readers' convenience, we have specialized the conditional sampler for such a noteworthy example in Section S6.3.

7. Illustrations. In this section, we display an illustrative example that provides evidence of the effectiveness of our proposal on simulated datasets. In Section S7, we also discuss how to adapt our strategy in presence of covariates and right-censored survival times, dealing with tumor survival data.

In the following simulated scenario, we have run the MCMC algorithm for 50,000 iterations with a burn-in period of 20,000 sweeps: the number of iterations allow the convergence of all the MCMC procedures. We have considered a synthetic dataset composed by $d = 3$ groups of observations, more specifically the data are generated by three different mixtures of Weibull distributions $f_1 \sim \text{Wbl}(1.5, 3/2)$, $f_2 \sim 0.5\text{Wbl}(2.5, 3/2) + 0.5\text{Wbl}(1.5, 3/2)$ and $f_3 \sim 0.5\text{Wbl}(2, 3/2) + 0.5\text{Wbl}(1.5, 3/2)$. For any $\ell = 1, 2, 3$, we have set $n_\ell = 100$. For the sake of simplifying notation, we denote by S_ℓ the survival function associated to f_ℓ, for any $\ell = 1, 2, 3$. As priors are concerned, we have used the specifications $a = a_0 = 1$, $b^{-1} = b_0^{-1} = 10$ and $T = 30$ which correspond to a noninformative specification, while we have fixed the values of $\sigma = \sigma_0 = 0.25$. We compare our BNP estimates (obtained via the marginal and conditional algorithm) with a frequentist estimate which is based on a smoothed version of the Kaplan–Meier estimator (using the optimized Matlab command ksdensity). Since the data have been generated independently across groups, we evaluate the frequentist estimator separately for each group. Figure 1 compares the true survival curve, the estimated
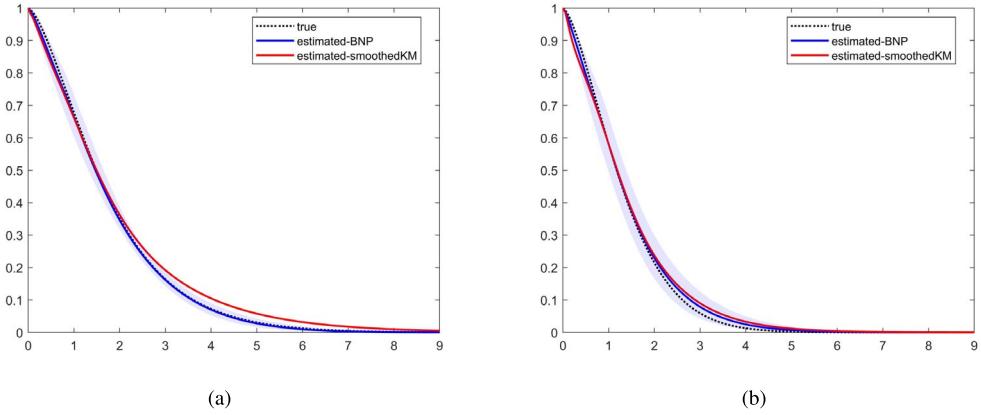
FIG. 1. *True survival functions (dotted) versus estimated (straight blue) for the first sample and 95% posterior credible intervals estimated using the marginal method (a) and conditional method (b). The BNP estimates are also compared with the corresponding frequentist estimate (straight red).*

survival function obtained via the marginal algorithm (in blue) and the frequentist estimate (in red). In Figure 1(a), the estimated credible intervals are obtained through the outputs of the marginal procedure, while in Figure 1(b) they are evaluated through the conditional method. More precisely, the BNP estimated survival function in Figure 1(a) has been obtained generating $N = 30{,}000$ values of $\mathbb{E}[\tilde{S}_\ell(t)|X, Y, T]$ (see Corollary 5) through the Gibbs sampler of Section 6.1, besides the respective credible intervals have been approximated evaluating the quantiles of the corresponding vector of outputs. On the other hand, as for Figure 1(b), we have implemented the conditional sampler depicted in Section 6.2 and we have estimated the posterior distribution (38) of $\tilde{S}_1$ to derive both the credible intervals and the estimator (posterior mean). Analogously Figures 2–3 compare the two survival functions for the second and third sample, respectively. First of all, we observe that both the BNP marginal and the BNP conditional method outperform with respect to the frequentist approach, providing estimates which are closer to the truth. This is also emphasized by the Kolmogorov distances $d_K(S_\ell, \tilde{S}_\ell)$ between the true survival curve and the estimated one reported in Table 1 for all $\ell = 1, 2, 3$ and for the different kinds of estimators. Second, it is apparent both marginal and conditional procedures yield similar point estimates, though the marginal method underestimates the credible intervals: indeed in such a situation the infinite dimensional random
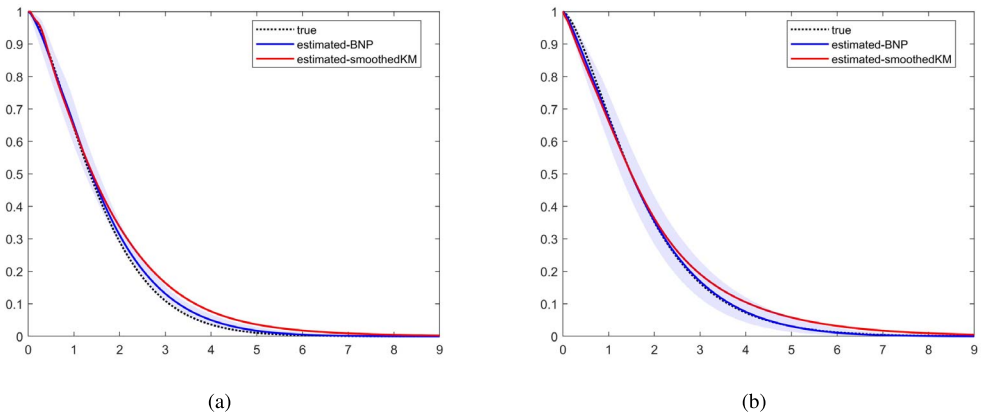

FIG. 2. *True survival functions (dotted) versus estimated (straight blue) for the second sample and 95% posterior credible intervals estimated using the marginal method (a) and conditional method (b). The BNP estimates are also compared with the corresponding frequentist estimate (straight red).*
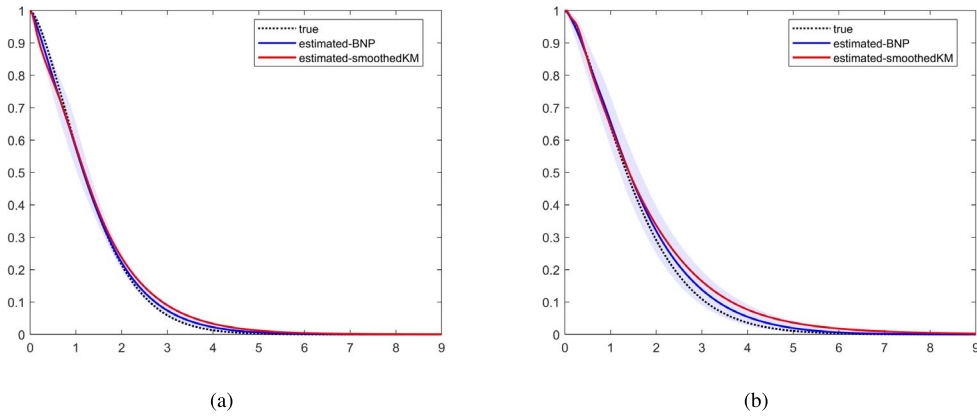
FIG. 3. *True survival functions (dotted) versus estimated (straight blue) for the third sample and 95% posterior credible intervals estimated using the marginal method (a) and conditional method (b). The BNP estimates are also compared with the corresponding frequentist estimate (straight red).*

TABLE 1
The Kolmogorov distance between the true and estimated survival functions for the different methodologies

	$d_K(S_1, \hat{S}_1)$	$d_K(S_2, \hat{S}_2)$	$d_K(S_3, \hat{S}_3)$
marginal BNP	0.033	0.027	0.023
conditional BNP	0.031	0.024	0.031
smoothed KM	0.060	0.042	0.056

elements are integrated out. We finally note that the conditional algorithm provides more reliable credible intervals which always contain the true survival functions, which is not the case for marginal sampling schemes (see Figure 3).

SUPPLEMENTARY MATERIAL

Supplement to "Survival analysis via hierarchically dependent mixture hazards" (DOI: 10.1214/20-AOS1982SUPP; .pdf). The supplementary material contains proofs, algorithmic details and additional illustrations.

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