

## Supplementary Material for Repulsive Mixtures

### 1 Supplementary concerning section 2.4

A possible issue with our repulsive mixture prior is that the full conditionals are nonstandard, complicating posterior computation. To address this, we propose a data augmentation scheme, introducing auxiliary slice variables to facilitate sampling [1]. In particular, letting  $g_0$  be a conjugate prior, introduce a latent variable  $u$  which is jointly modeled with  $\gamma$  through

$$\pi(\gamma_1, \dots, \gamma_k, u) \propto \left( \prod_{h=1}^k g_0(\gamma_h) \right) 1\{h(\gamma_1, \dots, \gamma_k) > u\}$$

Here  $1(A)$  is the indicator function, equalling 1 if the event  $A$  occurs, otherwise being 0. Marginalizing out  $u$ , we recover the original density  $\pi(\gamma_1, \dots, \gamma_k)$ . This algorithm is a slice sampling algorithm [4], a class of Markov chain Monte Carlo algorithms widely utilized for posterior inference in infinite mixture models [3]. To sample from 1, the derivation of  $A_j$  and therefore the invertibility of  $h$  is essential. For a repulsion function defined as (4), define  $A_j \equiv \bigcap_{\{s: s \neq j\}} [\gamma_j : g\{d(\gamma_s, \gamma_j)\} > u]$ . As long as  $g$  is invertible in its argument, the set  $A_j$  can be calculated, making sampling straightforward. When the repulsion function is defined as (3), one can introduce a latent variable for each product term. Finally, when  $m > 1$ , the location parameter vector can be sampled element-wise from the truncated distribution. Details on full conditional distributions can be found in the supplementary material. For simplicity, assume that  $h$  is defined as (4),  $g_0$  is the  $m$ -variate standard normal density and  $\phi$  is the  $m$ -variate spherical normal kernel. Let  $s_i = 1, \dots, k$  be the variable indicating which cluster the  $i$ th observation belongs to. Let  $n_j$  be the number of data points in the  $j$ th cluster and let  $\bar{y}_j$  be the average of observations in the  $j$ th cluster. Let  $\alpha_p = (\alpha_1 + n_1, \dots, \alpha_k + n_k)$ . Then the sampling algorithm can be summarized by the following steps:

*Step 1.* Update  $s_i$  for  $i = 1, \dots, n$  by multinomial sampling

$$(s_i | -) \sim \text{Multinomial}(l_1, \dots, l_k), \quad l_j = \frac{p_j \phi(y_i; \gamma_j, \sigma_j I)}{\sum_{h=1}^k p_h \phi(y_i; \gamma_h, \sigma_h I)};$$

*Step 2.* Sample  $(\gamma_j, \sigma_j)$  for  $j = 1, 2, \dots, k$  from

$$(\gamma_j | -) \sim N\left\{(1 + n_j/\sigma_j)^{-1} \bar{y}_j n_j / \sigma_j, I(1 + n_j/\sigma_j)^{-1}\right\} 1(\gamma_j \in A_j)$$

$$(1/\sigma_j | -) \sim \text{Ga}\left\{a_\sigma + \frac{n_j m}{2}, b_\sigma + \frac{1}{2} \sum_{\{i: s_i = j\}} (y_i - \gamma_j)^T (y_i - \gamma_j)\right\}$$

*Step 3.* Sample  $u$  and  $p$  from

$$(u | -) \sim \text{Un}\{0, h(\gamma)\}, \quad p \sim \text{Dirichlet}(\alpha_p)$$

### 2 Supplementary material concerning section 4

Let us provide more details about the densities in figure 2. Density (I) is a Students t density with eight degrees of freedom. Density (IIa) is a two-components mixture of Gaussians with mixture weights (0.3, 0.7), location parameters (−0.8, 0.8) and variances (0.2, 0.2). Density (IIb) is a mixture having the same weights and scale parameters as density (IIa) but location parameters (−1.5, 1.5), resulting in better separated clusters. Density (IIIa) is a mixture of a Gaussian with mean 0.7, variance 0.2 and weight 0.7 and a Pearson density with mean −0.7, variance 0.2, weight 0.3, skewness parameter −0.5 and kurtosis parameter 3. Density (IIIb) is a mixture having the same weights, scale parameters, skewness and kurtosis parameters as density (IIIa) but having location parameter (−1.2, 1.2), resulting in better separated clusters. Density (IV) is a bivariate mixture of two Gaussians with weight 0.5, location parameters (0, 0) and (2, 1), variances (0.2, 0.2) and (0.1, 0.1) and correlation coefficients 0.7 and 0.

Hyperparameters  $a_\sigma$  and  $b_\sigma$  for the density of the scale parameter were set to 3 and 1 respectively. Parameters  $\alpha_j$ s were all set equal to the same value  $\tilde{\alpha}$  and in accordance with [5]’s specification for

the density of the weights. For the non-repulsive model, the kernel locations were given independent standard normal priors. For the repulsive model, we considered a repulsion function defined as (4), with  $g$  defined as (5) and we chose  $g_0$  to be the standard normal. Concerning parameters involved in the repulsion function,  $\nu$  was set equal to 2 while  $\tau$  was chosen to guarantee a certain separation probability. In particular, the probability used to calibrate  $\tau$  was chosen to be 0.90 and  $c$  in definition 2 was fixed at 2.

The misclassification error was established in terms of divergence between the true similarity matrix and the posterior similarity matrix. Let  $n$  be the number of observations, then the similarity matrix is a  $(n \times n)$  matrix with  $(i, j)$  element equal to one if the  $j$ th and the  $i$ th observation belong to the same group and zero otherwise. As for the true similarity matrix, observations drawn from the same mixture component were considered as belonging to the same group. Let  $S$  be the true similarity matrix and  $\hat{S}_h$  be the similarity matrix obtained at the  $h$ th Markov chain Monte Carlo iteration. Let  $S(i, j)$  be the  $(i, j)$  element of  $S$  and define  $m_h$  as

$$m_h = \sum_{i=1}^n \sum_{j=i+1}^n 1(\hat{S}_h(i, j) \neq S(i, j))$$

with  $1(A)$  being the indicator function, equalling 1 if the event  $A$  occurs, otherwise being 0. The misclassification error at the  $h$ th iteration was calculated dividing  $m_h$  by the total number of pairs of  $n$  elements.

An approximation of the Kullback-Leibler divergence at the  $h$ th iteration was derived as

$$kl_h = \sum_{j=1}^s \log f_0(y_{0j}) / f(y_{0j}; \theta_h)$$

with  $f_0$  being the true density,  $f$  the fitted density,  $\theta_h$  the posterior sample at the  $h$ th iteration of parameters involved in  $f$  and  $y_{0j}$  for  $i = 1, \dots, s$  being  $s$  draws from the true density  $f_0$ . In all simulation examples  $s$  was set to 10,000.

As mentioned in section 4, knowing that the smoothing parameter  $\tilde{\alpha}$  directly affects the behavior of the mixture weights, it might be argued that under an accurate choice of  $\tilde{\alpha}$ , the non-repulsive prior may perform as well as the repulsive prior in emptying the extra components. Hence, we ran the non-repulsive model for different values of  $\tilde{\alpha}$ . This comparison was done by utilizing dataset (IIb) in figure 2. Table 1 provides posterior summary statistics for parameters involved in the repulsive model and non-repulsive model for different choices of  $\tilde{\alpha}$ . Clearly, as  $\tilde{\alpha}$  decreases, the non-repulsive model empties the extra components. However, we also see that the 95% credible interval of the location parameters now does not include the true value. This might be explained by the fact that as lower values of  $\tilde{\alpha}$  are considered, the posterior can concentrate on too few components leading to degenerate results in terms of estimates of specific component parameters.

As mentioned in section 2.4, parameter  $\tau$  in the repulsion function was calibrated to reach a particular separation probability. In order to assess the sensitivity of results to parameter's calibration, the KL divergence and the sum of extra weights were computed for different choices of separation probability. For this comparison, datasets (IIa) and (IIb) in figure 2 were considered. In practice, at each Markov chain Monte Carlo iteration, an approximation of the KL divergence and the sum of extra weights were obtained. Figure 1 and 2 show the median of these quantities over Markov chain Monte Carlo iterations. Clearly, as the probability of separation among clusters increases, the KL divergence between the fitted density and the true density increases while the sum of extra components decreases. However, as the sample size increase from  $n = 100$  to  $n = 1000$  the sensitivity of the two quantity of interest to the parameter's choice appears to vanish. Therefore, we expect that as the sample size increases, the choice of the separation probability will not excessively affect results.

### 3 Assumptions, cited theorems and proofs

#### 3.1 Assumptions

##### 3.1.1 Assumptions B1-B5

Assumptions B1-B5 corresponds to assumptions A1-A5 in [5]. Assumptions differ only in the conditions concerning the prior on the component-specific parameters in assumption A5. In condition

B5, we assume that  $\pi$  is defined as (2) and  $h$  is defined as either (3) or (4). For the sake of clarity, let us state assumption B1:

B1) There exists a  $q \geq 0$  such that for  $\delta_n = (\log n)^q n^{-1/2}$  the following holds

$$\lim_{M \rightarrow \infty} \limsup_{n \rightarrow \infty} E_n^0 \{ \Pi(\|f - f_0\|_1 \geq M\delta_n | Y_n) \} = 0$$

### 3.1.2 Conditions (i), (ii) and (iii) in theorem 3.1 of [6]

(i) The prior on  $\sigma$  has a continuous and positive Lebesgue density  $\psi$  on an interval containing  $\sigma_0$  and its distribution function  $\Psi$ , for constants  $e_1, e_2, e_3 > 0$ , satisfies

$$\Psi(s) \leq \exp(-e_1 s^{-e_2}) \text{ as } s \rightarrow 0 \text{ and } 1 - \Psi(s) \leq s^{-e_3} \text{ as } s \rightarrow \infty$$

(ii) The prior for the number of components is such that, for constants  $d_1, d_2 > 0$ ,

$$0 < \mu(k) \leq d_1 \exp(-d_2 k) \text{ for all } k \in \mathbb{N}$$

(iii) For each  $k$ , the prior for the weights is a Dirichlet with parameters  $(\alpha_1, \dots, \alpha_k)$  such that, for constants  $a_1, a_2 > 0, a_3 \geq 1$  and for  $0 < \epsilon \leq 1/(a_3 k)$  and  $j = 1, \dots, k$

$$a_2 \epsilon^{a_1} \leq \alpha_j \leq a_3$$

### 3.2 Cited theorems

**Theorem 1** ([2]). *Let  $\pi_n$  be a sequence of priors on a class of densities  $\mathcal{F}$  equipped with a metric  $d$  that can be either the Hellinger or the one induced by the  $L_1$ -norm. Assume that for positive sequences  $\bar{\epsilon}_n, \tilde{\epsilon}_n \rightarrow 0$  such that  $n \min(\bar{\epsilon}_n, \tilde{\epsilon}_n) \rightarrow \infty$ , constants  $d_1, d_2, d_3, d_4 > 0$  and sets  $\mathcal{F}_n \subseteq \mathcal{F}$ , we have*

$$\log D(\bar{\epsilon}_n, \mathcal{F}_n, d) \leq d_1 n \bar{\epsilon}_n^2 \quad (1)$$

$$\pi_n(\mathcal{F} \setminus \mathcal{F}_n) \leq d_3 \exp\{-(d_2 + 4)n\tilde{\epsilon}_n^2\} \quad (2)$$

$$\pi_n\{B_{KL}(f_0; \tilde{\epsilon}_n^2)\} \geq d_4 \exp(-d_2 n \tilde{\epsilon}_n^2) \quad (3)$$

where  $B_{KL}(f_0; \tilde{\epsilon}_n^2) = \{f : \int f_0 \log(f_0/f) \leq \tilde{\epsilon}_n^2; \int f_0 \log(f_0/f)^2 \leq \tilde{\epsilon}_n^2\}$ .

Then, for  $\epsilon_n = \max(\bar{\epsilon}_n, \tilde{\epsilon}_n)$  and a sufficiently large constant  $M > 0$ , the posterior probability

$$\pi_n\{f : d(f, f_0) > M\epsilon_n | Y_n\} \rightarrow 0$$

in  $P_0^n$  probability, as  $n \rightarrow \infty$ .

### 3.3 Proofs

*Proof of lemma 1.* By assumption B0,  $\mu(k = k_0) > 0$ . We consider the case  $f$  is a finite mixture with  $k_0$  components. By assumption A1, for each  $\eta > 0$  there is a corresponding  $\delta > 0$  such that, for any given  $y \in \mathcal{Y}$  and for all  $\gamma_1, \gamma_2 \in \Gamma$  with  $|\gamma_1 - \gamma_2| < \delta$ , we have that  $|\phi(y; \gamma_1) - \phi(y; \gamma_2)| < \eta$ . Let  $S_\delta = \Gamma_\delta \times P_\delta$  with  $\Gamma_\delta = \{\gamma : |\gamma_j - \gamma_{0j}| \leq \delta, j \leq k_0\}$  and  $P_\delta = \{p : |p_j - p_{0j}| \leq \delta, j \leq k_0\}$ . By assumption A1 and A2, for any given  $y$  and for any  $\eta > 0$ , there is a  $\delta > 0$  such that  $|f_0 - f| \leq \eta$  if  $\theta \in S_\delta$ . This means that,  $f \rightarrow f_0$  as  $\theta \rightarrow \theta_0$ , for any given  $y$ . Equivalently, we can say that  $|\log(f_0/f)| \rightarrow 0$  pointwise as  $\theta \rightarrow \theta_0$ . Notice that

$$|\log(f_0/f)| \leq \left| \log \left\{ \sup_{\gamma \in D_0} \phi(\gamma) \right\} - \log \left\{ \inf_{\gamma \in D_0} \phi(\gamma) \right\} \right|$$

By assumption A3 and applying the dominated convergence theorem, for any  $\epsilon > 0$  there is a  $\delta > 0$  such that  $\int f_0 \log(f_0/f) < \epsilon$  if  $\theta \in S_\delta$ . By the independence of the weights and the parameters of the kernel,

$$\Pi(KL(f_0, f) < \epsilon) \geq \lambda(P_\delta) \pi(\Gamma_\delta)$$

Assumption A4 combined with the fact that  $\{\gamma : \|\gamma - \gamma_0\|_1 \leq \delta\} \subseteq \Gamma_\delta$  result in  $\pi(\Gamma_\delta) > 0$ . Finally, since  $\lambda = \text{Dirichlet}(\alpha)$ , it can be shown that  $\lambda(P_\delta) > 0$ .  $\square$

*Proof of lemma 2.* Let  $D = \{\gamma : \|\gamma - x\|_1 < v/2\}$ . By the assumptions on  $h$ , given a vector  $x$  satisfying condition A4 in lemma 2,  $h(\gamma) > 0$  for  $\gamma$  such that  $d(\gamma_s, x_s) < v/2$  for  $s = 1, \dots, k$ . Since,

$$D \subseteq \{\gamma : d(\gamma_s, x_s) < v/2; s = 1, \dots, k\},$$

it follows that  $h(\gamma) > 0$  on  $D$ . By assumption,  $g_0$  is positive on  $\Gamma$ , therefore it follows that  $\pi(\gamma) > 0$  on  $D$ .  $\square$

*Proof of lemma 3.* To prove lemma 3 we need to show that the three conditions of theorem 2.1 in [2] are satisfied. First, define  $D(\epsilon, \mathcal{F}, d_s)$  as the maximum number of points in  $\mathcal{F}$  such that the distance, with respect to metric  $d_s$ , between each pair is at least  $\epsilon$ . Let  $d_s$  be either the Hellinger metric or the one induced by the L1-norm. For given sequences  $k_n, a_n, u_n \uparrow \infty$  and  $b_n \downarrow 0$  define

$$\mathcal{F}_n^{(k)} = \left\{ f : f = \sum_{j=1}^k p_j \phi(\gamma_j, \sigma), \gamma_j \in (-a_n, a_n)^k, \sigma \in (b_n, u_n) \right\}$$

and  $\mathcal{F}_n = \cup_{j=1}^{k_n} \mathcal{F}_n^{(j)}$ . As it is shown in [6], for constants  $f_2 \geq f_1 > 0$  and  $l_1, l_2, l_3 > 0$ , derived below to satisfy condition (2) and (3) in [2], and defining  $f_1 \log n \leq k_n \leq f_2 \log n$ ,  $a_n = l_3 (\log \bar{\epsilon}_n^{-1})^{1/2}$ ,  $b_n = l_1 (\log \bar{\epsilon}_n^{-1})^{-1/e_2}$  and  $u_n = \bar{\epsilon}_n^{-l_2}$ ,  $\log D(\bar{\epsilon}_n, \mathcal{F}_n, d_s) \lesssim n \bar{\epsilon}_n^2$  with  $\bar{\epsilon}_n = n^{-1/2} \log n$ .

Let  $A_{n,j} = (-a_n, a_n)^j$ . In order to show condition (2) of theorem 2.1. in [2], we need to show that there is a constant  $q_1 > 0$  such that  $\pi(A_{n,k}^C) \lesssim \exp(-q_1 a_n^2)$ . From the exchangeability assumption it follows

$$\begin{aligned} pr(A_{n,k}^C | k = s) &= \sum_{j=1}^s \frac{s!}{j!(s-j)!} \pi(A_{n,j}^C \times A_{n,s-j}) \\ &\leq s \sum_{j=1}^s \frac{(s-1)!}{(j-1)!(s-j)!} \pi(A_{n,j}^C \times A_{n,s-j}) \leq s \pi_m(A_{n,1}^C) \end{aligned}$$

Therefore, condition C1 implies that, for a positive constant  $q_1$  we have  $\pi(A_{n,k}^C) \lesssim E(k) \exp(-q_1 a_n^2)$  with  $E(k) < \infty$  by condition (ii). Given a positive constant  $z_2$  chosen to satisfy condition (3) in theorem 2.1 of [2], let  $f_1 \geq (z_2 + 4)/d_2$ ,  $l_1 \leq \{e_1/4(z_2 + 4)\}^{1/e_2}$ ,  $l_2 \geq 4(z_2 + 4)/e_3$  and  $l_3 \geq \{4(z_2 + 4)/q_1\}^{1/2}$ . Under these values of  $f_1, l_1, l_2$  and  $l_3$ , following [6], assumptions (i), (ii) and assumption C1 imply  $\Pi(\mathcal{F} \setminus \mathcal{F}_n) \lesssim \exp\{-(z_2 + 4)n \bar{\epsilon}_n^2\}$  with  $\bar{\epsilon}_n = n^{-1/2}(\log n)^{1/2}$ .

To show condition (3) of theorem 2.1 in [2], we can again follow the proof of theorem 3.1. in [6]. The only thing we need to show is that, there are constants  $u_1, u_2, u_3 > 0$  such that for any  $\epsilon_n \leq u_3$

$$\pi(\|\gamma - \gamma_0\|_1 \leq \epsilon_n) \geq u_1 \exp\{-u_2 k_0 \log(1/\epsilon_n)\}$$

that is guaranteed by condition C2. Therefore, it can be easily showed that, for sufficiently large  $n$ ,  $z_2 > 0$  and  $\bar{\epsilon}_n = n^{-1/2}(\log n)^{1/2}$ ,  $\Pi\{B_{KL}(f_0, \bar{\epsilon}_n^2)\} \gtrsim \exp(-z_2 n \bar{\epsilon}_n^2)$ .  $\square$

*Proof of lemma 4.* First, let us check that condition C1 is satisfied. Clearly, under the assumptions on  $h$ ,  $\pi$  leads to exchangeable atoms. Under the assumptions on  $\pi$ , the following holds

$$\pi_m(|\gamma_1| \geq t) = \int_{|\gamma_1| \geq t} \pi_m(\gamma_1) d\gamma_1 \leq c_1 c_2 \int_{|\gamma_1| \geq t} g_0(\gamma_1) d\gamma_1$$

with  $c_1$  and  $c_2$  defined as in (2). It follows that there exists a constant  $n_1 > 0$  such that  $\pi_m(|\gamma_1| \geq t) \lesssim \exp(-n_1 t^2)$ .

Now let us verify condition C2. Assumptions on  $h$  imply that for any  $0 < \epsilon < 1$  there is a corresponding  $0 < \delta = g^{-1}(\epsilon)$  and constants  $w_1 > 0$  such that  $h(\gamma) \geq w_1 \epsilon^{k_0}$  for all  $\gamma$  satisfying  $\min_{\{(s,j): s < j\}} d(\gamma_j, \gamma_s) \geq \delta$ . Let  $u_3$  be defined as

$$u_3 = \min[\epsilon_1/2, g(\delta_1)]$$

with  $\epsilon_1$  defined as in assumption B0 and  $\delta_1 = \epsilon_1(1 - 1/k_0)$ . By assumption  $\epsilon < u_3$  and therefore  $\delta < \delta_1$ . Let us define  $M(\gamma, x)$  and  $N(\gamma, x)$  as follows,

$$M(\gamma, x) = \left\{ \gamma : \min_{\{(s,j): s < j\}} d(\gamma_j, \gamma_s) \geq x \right\}, \quad N(\gamma, x) = \{\gamma : |\gamma_j - \gamma_{0j}| \leq x; j = 1, \dots, k_0\}$$

Then,

$$\begin{aligned}
\pi(\|\gamma - \gamma_0\|_1 \leq \epsilon) &\geq \int_{\{\|\gamma - \gamma_0\|_1 \leq \epsilon\} \cap M(\gamma, \delta)} \pi(\gamma) d\gamma \\
&\gtrsim \int_{\{\|\gamma - \gamma_0\|_1 \leq \epsilon\} \cap M(\gamma, \delta)} \epsilon^{k_0} \prod_{j=1}^{k_0} g_0(\gamma_j) d\gamma \\
&\gtrsim \int_{N(\gamma, \epsilon/k_0) \cap M(\gamma, \delta_1)} \epsilon^{k_0} \prod_{j=1}^{k_0} g_0(\gamma_j) d\gamma
\end{aligned}$$

Now let us show that  $N(\gamma, \epsilon/k_0) \subseteq M(\gamma, \delta_1)$ . Consider the pair  $(s, j)$  with  $s \neq j$ . Without loss of generality assume  $\gamma_{0s} > \gamma_{0j}$ . Now, consider the possible values of  $(\gamma_j, \gamma_s)$  contained in the set  $N(\gamma, \epsilon/k_0)$ . The smallest distance between values of  $\gamma_s$  and  $\gamma_j$  contained in  $N(\gamma, \epsilon/k_0)$  is

$$(\gamma_{0s} - \epsilon/k_0) - (\gamma_{0j} + \epsilon/k_0) \geq \epsilon_1 - 2\epsilon/k_0 \geq \epsilon_1(1 - 1/k_0) = \delta_1$$

Since the previous holds for any pair  $(s, j)$  we have  $N(\gamma, \epsilon/k_0) \subseteq M(\gamma, \delta_1)$ . Therefore,

$$\begin{aligned}
\pi(\|\gamma - \gamma_0\|_1 \leq \epsilon) &\gtrsim \int_{N(\gamma, \epsilon/k_0)} \epsilon^{k_0} \prod_{j=1}^{k_0} g_0(\gamma_j) d\gamma \\
&\gtrsim \epsilon^{k_0} \exp\{-g_1 k_0 \log(1/\epsilon)\} \\
&\gtrsim \exp\{-(g_1 + 1)k_0 \log(1/\epsilon)\}
\end{aligned}$$

for a constant  $g_1 > 0$ . □

*Proof of theorem 1.* Only for this proof and for ease of notation the density  $f$  will be referred as  $f_\theta$ . Define the non identifiability set as  $T = \{\theta : f_\theta = f_0\}$ . In order to define each vector in  $T$ , let  $0 = t_0 < t_1 < t_2 \dots < t_{k_0} \leq k$  and  $\gamma_j = \gamma_{0i}$  for  $j \in I_i = \{t_{i-1} + 1, t_i\}$ . Let  $p_{0i} = \sum_{j=t_{i-1}+1}^{t_i} p_j$  and  $p_j = 0$  for  $j > t_{k_0}$ . Define  $q_j = p_j/p_{0i}$  for  $j \in I_i$ . Define  $A_n = \left\{ \min_{\sigma \in S_k} \left( \sum_{i=1}^{k-k_0} p_{\sigma(i)} \right) > \delta_n M_n \right\}$  and  $A'_n = A_n \cap \{\|f - f_0\|_1 \leq M\delta_n\}$ . Let  $D_n = \int_{\{\|f - f_0\|_1 < \delta_n\}} \exp(l_n(\theta) - l_n(\theta_0)) d(\pi \times \lambda)(\theta)$  with  $l_n(\theta_0)$  being the log-likelihood evaluated at  $\theta_0$ . Along the line of [5]'s proof, to prove theorem 1 we need to show that for any  $\epsilon > 0$  there are positive constants  $m_1, m_2$  and a permutation  $\sigma \in S_k$  such that

$$D_n \geq m_1 n^{-s(k_0, \alpha)/2} \quad (4)$$

$$\Pi(A'_n) \leq m_2 \delta_n^{s(k_0, \alpha)} M_n^{\bar{\alpha} - m/2 - r_2} \quad (5)$$

with  $s(k_0, \alpha) = k_0 - 1 + mk_0 + \sum_{j=1}^{k-k_0} \alpha_{\sigma(j)}$ . Following [5]'s proof, we can show that, under condition B5, (4) is satisfied for sufficiently large  $n$ . Concerning (5), [5] showed that on  $A'_n$ , there is a set  $I_i$  containing indices  $j_1$  and  $j_2$  such that

$$|\gamma_{j_1} - \gamma_{0i}| \leq (\delta_n/q_{j_1})^{1/2}, \quad |\gamma_{j_2} - \gamma_{0i}| \leq (\delta_n/q_{j_2})^{1/2}$$

with  $q_{j_1} > \epsilon/k_0$  and  $q_{j_2} > \delta_n M_n/2$ . The triangle inequality implies

$$|\gamma_{j_1} - \gamma_{j_2}| \leq 2 \{\delta_n / \min(q_{j_1}, q_{j_2})\}^{1/2}$$

Now, for sufficiently large  $n$ ,  $\min(q_{j_1}, q_{j_2}) > \delta_n M_n/2$  and therefore  $|\gamma_{j_1} - \gamma_{j_2}| \lesssim M_n^{-1/2}$ . Since  $g$  is bounded above by a positive constant, it exists a constant  $c > 0$  such that

$$h(\gamma) \leq cg \{d(\gamma_{j_1}, \gamma_{j_2})\} \leq cg \left( M_n^{-1/2} \right) \quad (6)$$

for  $\gamma \in A'$ . Let the prior probability of the set  $A'_n$  be defined as  $\Pi(A'_n) = \int_{A'_n} d(\pi \times \lambda)(\gamma \times p)$ . To find an upper bound for this integral, directly apply the proof of [5] showing that  $\Pi(A'_n) \leq g \left( M_n^{-1/2} \right) \delta_n^{s(k_0, \alpha)} M_n^{\bar{\alpha} - m/2}$ . By assumption, for sufficiently large  $n$ ,  $g \left( M_n^{-1/2} \right) \leq r_1 M_n^{-r_2}$ . Letting  $s_{r_2} = r_2 + m/2 - \bar{\alpha}$ , it follows

$$\Pi(A'_n) \leq M_n^{-s_{r_2}} \delta_n^{s(k_0, \alpha)}$$

□

## References

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Table 1: Percentiles 2.5th, 50th and 97.5th of location parameters involved in the two components with highest weights and sum of extra weights under repulsive atoms and non-repulsive atoms for different values of  $\tilde{\alpha}$  under dataset (*IIb*)

$\tilde{\alpha}$	n=100				n=1000			
	Non-Repulsive			Repulsive	Non-Repulsive			Repulsive
	1/3	1/10	1/100	1/3	1/3	1/10	1/100	1/3
Parameter 1 (truth 1.50)								
2.5	1.59	1.61	1.61	1.53	1.51	1.52	1.52	1.49
50	1.62	1.61	1.61	1.64	1.52	1.52	1.52	1.53
97.5	1.68	1.64	1.63	1.75	1.54	1.53	1.52	1.56
Parameter 2 (truth -1.50)								
2.5	-1.50	-1.48	-1.42	-1.70	-1.51	-1.50	-1.49	-1.55
50	-1.39	-1.39	-1.39	-1.52	-1.48	-1.48	-1.48	-1.49
97.5	-1.21	-1.28	-1.32	-1.34	-1.44	-1.47	-1.47	-1.44
Extra weights sum ( $\times 10^3$ )								
2.5	4	0	0	1	1	0	0	0
50	69	8	1	10	12	1	1	1
97.5	277	139	15	43	138	25	4	4

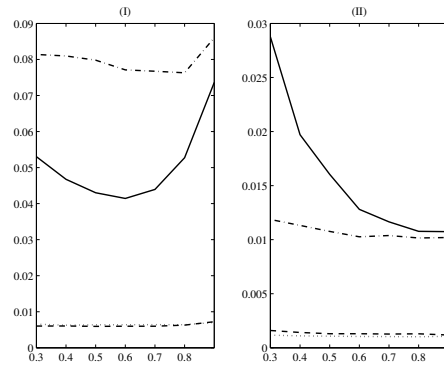


Figure 1: Plot of KL divergence (*I*) and sum of extra weights (*II*) for different choice of separation probability (x axis) under dataset (*IIa*) for  $n = 100$  (solid) and  $n = 1000$  (dash) and dataset (*IIb*) for  $n = 100$  (dash-dot) and  $n = 1000$  (dot) under a mixture of six components

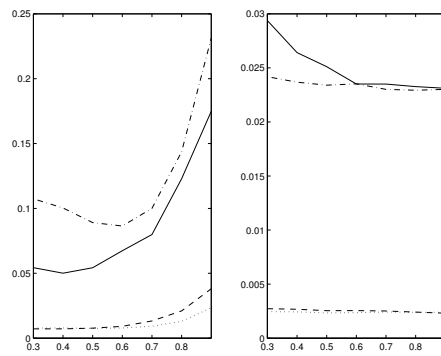


Figure 2: Plot of KL divergence ( $I$ ) and sum of extra weights ( $II$ ) for different choice of separation probability (x axis) under dataset ( $IIa$ ) for  $n = 100$  (solid) and  $n = 1000$  (dash) and dataset ( $IIb$ ) for  $n = 100$  (dash-dot) and  $n = 1000$  (dot) under a mixture of ten components