

Computational methods for Bayesian nonparametric models

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Structure

- Pólya urn scheme samplers – Conjugate/non-conjugate DPMM, split-and-merge.
- Stick-breaking – Block Gibbs sampler, slice sampler
- Normalized random measures – Simulation, Truncation methods, slice sampling

Not covered

- Feature allocation – Indian buffet process, beta-Bernoulli processes and generalisations
- More complicated processes – Poisson-Kingman processes, etc.
- Distributed computing
- Sequential Monte Carlo
- Variational Bayes
- More complicated models – hierarchical processes, nested processes, dependent processes, etc.

Bayesian nonparametric mixture models

Hierarchical model

$$y_i | \theta_i \sim k(y_i | \theta_i), \quad \theta_i | F \sim F, \quad F \sim \text{DP}(M, H)$$

Allocation representation

$$y_i | s_i, \theta_1^*, \theta_2^*, \dots \sim k(y_i | \theta_{s_i}^*),$$

$$p(s_i = k) = w_k \sim \text{GEM}(M)^1, \quad \theta_k^* \overset{i.i.d.}{\sim} H$$

where $\theta_1^*, \theta_2^*, \dots$ are the **distinct values**

¹Griffiths-Engen-McCloskey process

Marginal MCMC methods for Dirichlet process mixture models

The key idea here is that the predictive distribution of the allocation variable s_j is described by a Chinese restaurant process

$$p(s_j = k | s_1, \dots, s_{n-1}) = \begin{cases} \frac{n_{k,j}}{M+j-1} & 1 \leq k \leq K_{j-1} \\ \frac{M}{M+j-1} & k = K_{j-1} + 1 \end{cases}$$

where K_{j-1} is the number of different values of s_1, \dots, s_{j-1} and $n_{k,j}$ is the number of observations in the k -th cluster.

Then

$$p(s_1, \dots, s_n) = \prod_{j=1}^n p(s_j | s_1, \dots, s_{j-1})$$

Marginal MCMC methods for Dirichlet process mixture models

The posterior is finite (but not fixed) dimensional

$$p(s_1, \dots, s_n, \theta_1^*, \dots, \theta_{K_n}^* | y) \propto \prod_{k=1}^{K_n} \prod_{\{i | s_i = k\}} k(y_i | \theta_k^*) \prod_{k=1}^{K_n} h(\theta_k^*) p(s_1, \dots, s_n)$$

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In a conjugate DPM we assume that $\int \prod_{\{i | s_i = k\}} k(y_i | \theta) h(\theta) d\theta$ can be calculate analytically.

$$p(s_1, \dots, s_n | y) = \int p(s_1, \dots, s_n, \theta_1^*, \dots, \theta_{K_n}^* | y) d\theta_1^* \dots d\theta_{K_n}^*$$

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Then

$$\begin{aligned} p(s_1, \dots, s_n | y) &= \int p(s_1, \dots, s_n, \theta_1^*, \dots, \theta_{K_n}^* | y) d\theta_1^* \dots d\theta_{K_n}^* \\ &\propto \prod_{k=1}^{K_n} \int \prod_{\{i | s_i = k\}} k(y_i | \theta_k^*) h(\theta_k^*) d\theta_k^* \times p(s_1, \dots, s_n) \end{aligned}$$

Gibbs sampler

Due to exchangeability

$$p(s_j = k | s_1, \dots, s_{n-1}) = \begin{cases} \frac{n_k^{-j}}{M+n-1} k(y_j | \{y_i | s_i = k\}) & 1 \leq k \leq K^{-j} \\ \frac{M}{M+n-1} k(y_j) & k = K^{-j} + 1 \end{cases}$$

where

- $n_k^{-j} = \#\{s_i = k, i \neq j\}$.
- $k(y_j | \{y_i | s_i = k\})$ is the predictive distribution with likelihood k and prior H
- K^{-j} is the number of distinct values in $s_1, \dots, s_{j-1}, s_{j+1}, \dots, s_n$.

Non-conjugate Dirichlet process mixtures – Neal's algorithm 8

We work on s_1, \dots, s_n and $\theta_1^*, \dots, \theta_{K_n}^*$. The posterior is proportional to

$$p(s_1, \dots, s_n) \prod_{i=1}^n k(y_i | \theta_{s_i}^*) \prod_{k=1}^{K_n} h(\theta_k^*)$$

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where

$$p(s_1, \dots, s_n) = M^{K_n} \frac{\Gamma(n)}{\Gamma(M+n)} \prod_{k=1}^{K_n} \Gamma(n_k)$$

and $n_k = \#\{s_i = k\}$. This is the **exchangeable product partition formula (EPPF)**.

Neal's algorithm 8

In this sampler, **latent variables** ψ_1, \dots, ψ_m are introduced to define the augmented posterior (by exchangeability, we only look at s_n).

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$$M^{K-n} \prod_{k=1}^m \left(\frac{M}{m} \right)^{I(s_n=K_{-n}+1 \text{ and } \theta_{K_{-n}+1}^*=\psi_k)} \\ \times \frac{\Gamma(n)}{\Gamma(M+n)} \prod_{k=1}^{K_n} \Gamma(n_k) \prod_{i=1}^n k(y_i | \theta_{s_i}^*) \prod_{k=1}^{K_{-n}} h(\theta_k^*) \prod_{k=1}^m h(\psi_k)$$

Full conditional distribution

To update the allocation variable for the i -th observations re-label the s 's so that $s_i \geq s_j$ for $1 \leq j \leq n$.

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$$p(s_i = k) \propto \begin{cases} n_{i,k} k(y_i | \theta_k^*) & 1 \leq k \leq K_{-n} \\ \frac{M}{m} k(y_i | \psi_{k-K_{-n}}) & K_{-n} + 1 \leq k \leq K_{-n} + m \end{cases}$$

where

- If the i -th observation is a singleton (i.e. $n_{s_i} = 1$) then $\psi_1 = \theta_{K_{-n}+1}^*$ and $\psi_2, \dots, \psi_m \sim H$.
- Otherwise, $\psi_1, \dots, \psi_m \sim H$.

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- Otherwise, $\psi_1, \dots, \psi_m \sim H$.

If $k > K_{-n}$, set $s_n = K_{-n} + 1$ and $\theta_{K_{-n}+1}^* = \psi_{k-K_{-n}}$.

Comments

- Algorithm 8 has been extended to the case of normalized random measures with independent increments by Favaro and Teh (2013).
- The approach can be interpreted as akin to a pseudo-marginal method (Andrieu and Roberts, 2009).

Split-merge samplers

Split-merge samplers try to make larger moves in allocation space.

Two approaches SAMS (Dahl and Newcomb, 2022) and RGMS (Neal and Jian, 2004) which differ in re-allocation mechanism.

Both algorithms involve sampling two observations i and j at random.

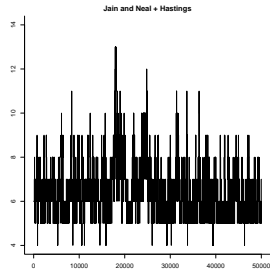
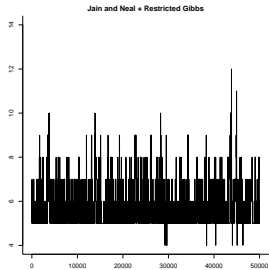
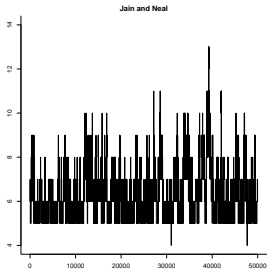
- If these $s_i \neq s_j$, merge.
- If there $s_i = s_j$, split. Create two clusters containing observation i and j respectively.
 - SAMS – Allocate all other observations with $s_k = s_i$ i one of the two new clusters using the full conditional (sequentially).
 - RGMS – Randomly allocate all observations with $s_k = s_i$ to the two clusters. Update the allocation using the full conditional distribution.

Joint work with Alessandro Colombi

These can be seen as a type of informed neighbourhood scheme where

- The neighbourhood is all possible re-clusterings of the two clusters conditional on the anchor points.
- The sequential/restricted Gibbs scan is a type of informed proposal.

We can build an adaptive random neighbourhood informed (ARNI) scheme where the random neighbourhood is a core-periphery model with a “score” of “core-ness” for each observation.



Conditional samplers for DPM models

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Usually, these algorithms work for both conjugate and non-conjugate DPM models.

Representations of the Dirichlet process

Suppose that $F \sim \text{DP}(M, H)$ then

$$F = \sum_{k=1}^{\infty} w_k \delta_{\theta_k}$$

where $\theta_k \sim H$ and ...

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2. Normalized random measure with independent increment

$w_k = \frac{\gamma_k}{\sum_{m=1}^{\infty} \gamma_m}$ where $\gamma_1, \gamma_2, \dots$ are the jumps of a gamma process.

Samplers using stick-breaking representations

Choose a value K and set $V_{K+1} = 1$ then only w_1, \dots, w_{K+1} are non-zero.

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The truncation error can be measured in the following way

$$\|f(y) - f_K(y)\|_1 < 4 \left(1 - \mathbb{E} \left[\left(\sum_{k=1}^K w_k \right)^n \right] \right)$$

where

- $f(y)$ is the joint predictive distribution of the data
- $f_K(y)$ is the joint predictive distribution of the data under the truncation.
- $\|\cdot\|_1$ represents L_1 distance.

Blocked Gibbs sampler (Ishwaran and James, 2001)

If we truncate at K components, then

$$p(s_1, s_2, \dots, s_n) = \prod_{i=1}^n w_{s_i} = \prod_{k=1}^K V_k^{n_k} (1 - V_k)^{m_k}$$

where $n_k = \#\{i | s_i = k\}$ and $m_k = \sum_{i=k+1}^K n_i$.

Blocked Gibbs sampler

1. The full conditional for s_i is

$$p(s_i = k) \propto w_k k(y_i | \theta_k), \quad k = 1, \dots, K + 1$$

2. The full conditional for V_k is $\text{Be}(1 + n_k, M + m_k)$.
3. The full conditional for θ_k is proportional to

$$h(\theta_k) \prod_{\{i | s_i = k\}} k(y_i | \theta_k)$$

Note: This full conditional distribution will have a known if the prior distribution of θ_k is conjugate.

Comments

- This method can be directly applied to the Pitman-Yor process where $V_k \sim \text{Be}(1 - a, M + k a)$.
- The form $p(s_1, s_2, \dots, s_n) = \prod_{k=1}^K V_k^{n_k} (1 - V_k)^{m_k}$ is convenient for other models such as probit stick-breaking (Rodriguez and Dunson, 2011)

A slice sampler for DPM models

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The **slice sampler** uses a random truncation to avoid a truncation error.

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$$p(y_i, u_i) = \sum_{k=1}^{\infty} \mathbb{I}(u_i < w_k) k(y_i|\theta_k)$$

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Slice sampler for DPM models

$$p(y_i, u_i, s_i) = \mathbb{I}(u_i < w_{s_i}) k(y_i | \theta_{s_i})$$

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Note: The marginal of y_i is unchanged.

The posterior distribution is

$$p(u, s, V, \theta | y) \propto h(\theta) p(V) \prod_{i=1}^n \mathbb{I}(u_i < w_{s_i}) k(y_i | \theta_{s_i})$$

Gibbs sampler (Walker, 2007)

1. Full conditional distribution of s_i

$$p(s_i = k) \propto I(w_k > u_i) k(y_i | \theta_k)$$

Note: there are only a finite number of non-zero values.

2. Full conditional of θ_k is

$$h(\theta_k) \prod_{\{i | s_i = k\}} k(y_i | \theta_k)$$

3. Full conditional distribution of $u_i \sim U(0, w_{s_i})$.
4. Full conditional distribution of V_k is proportional to

$$p(V_k) \prod_{i=1}^n I \left(u_i < V_{s_i} \prod_{j < s_i} (1 - V_j) \right).$$

Accelerated Gibbs sampler (Kalli et al., 2011)

It is more efficient to jointly updating v_1, \dots, v_K and u_1, \dots, u_n using

$$p(v_1, \dots, v_K, u_1, \dots, u_n) = p(u_1, \dots, u_n | v_1, \dots, v_K) p(v_1, \dots, v_K)$$

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$$p(v_1, \dots, v_K, u_1, \dots, u_n) = p(u_1, \dots, u_n | v_1, \dots, v_K) p(v_1, \dots, v_K)$$

Then steps 3 and 4 become

1. The full conditional for V_k is $\text{Be}(1 + n_k, M + m_k)$. *i.e.* full conditional from blocked Gibbs sampler.
2. Full conditional distribution of $u_i \sim \text{U}(0, w_{S_i})$.

To complete step 1, we only need $w_1, \dots, w_K > \min\{u_i\}$.

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Notice that $w_k < \prod_{j=1}^K (1 - V_j)$ for $k > K$ and that, for $k > \max\{s_i\}$, $V_k \sim \text{Be}(1, M)$.

To complete step 1, we only need $w_1, \dots, w_K > \min\{u_i\}$.

Notice that $w_k < \prod_{j=1}^K (1 - V_j)$ for $k > K$ and that, for $k > \max\{s_i\}$, $V_k \sim \text{Be}(1, M)$.

So we choose the first K for which $\prod_{j=1}^K (1 - V_j) < \min\{u_i\}$ which implies that $w_k < \min\{u_i\}$ for $k > K$.

Completely random measures

$\tilde{\mu}$ is a **completely random measure** (CRM) on \mathbb{X} if, for any disjoint subsets A_1, \dots, A_n , $\tilde{\mu}(A_1), \dots, \tilde{\mu}(A_n)$ are mutually independent.

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We concentrate on **completely random measures** (CRM's) which can be represented in terms of **jump sizes** J_i and **jump locations** X_i as

$$\tilde{\mu} = \sum_{i=1}^{\infty} J_i \delta_{X_i}$$

where δ is Dirac's delta function and have Lévy-Khintchine representation

$$\mathbb{E} \left[e^{-\int f(x) \tilde{\mu}(dx)} \right] = e^{-\int_0^\infty \int [1 - e^{-sf(x)}] h(dx) \nu(ds)}$$

where $h(\cdot)$ is a p.d.f. and $\nu(\cdot)$ is a Lévy intensity

$$\int_{\mathbb{R} \setminus \{0\}} \min(1, x^2) \nu(x) dx < \infty.$$

Simulating completely random measures

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Simulating completely random measures

Suppose that we wish to simulate a realisation of a Lévy process with intensity ν .

Define the **tail mass function** η of a Lévy process with Lévy intensity ν to be $\eta(x) = \int_x^\infty \nu(z) dz$.

Ferguson and Klass (1972) showed the Lévy process can be represented as

$$F = \sum_{k=1}^{\infty} J_k \delta_{\theta_k}$$

where $\gamma_k = \eta^{-1}(E_j)$ and E_1, E_2, \dots are the points of a Poisson process with intensity 1.

Example: Gamma process

Recall that a gamma process has Lévy intensity
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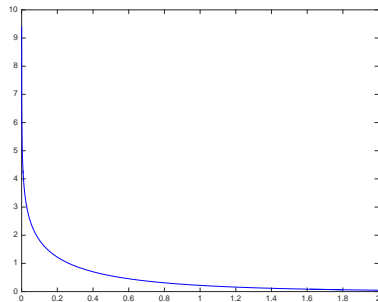
The tail mass integral is

$$\eta(x) = M \int_x^\infty z^{-1} \exp\{-z\} dz = M \operatorname{Ei}(x)$$

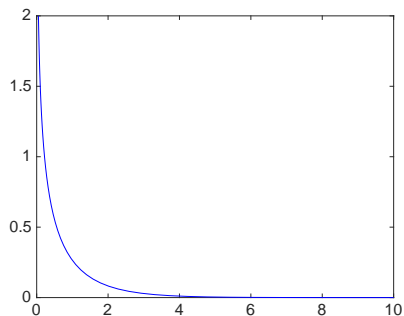
where $\operatorname{Ei}(x)$ is the exponential-integral function.

Gamma process simulation

$$\eta(x)$$

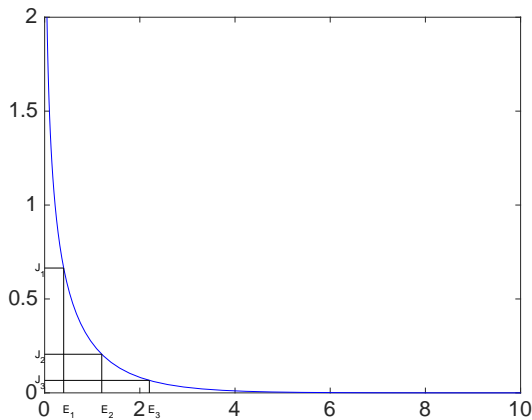


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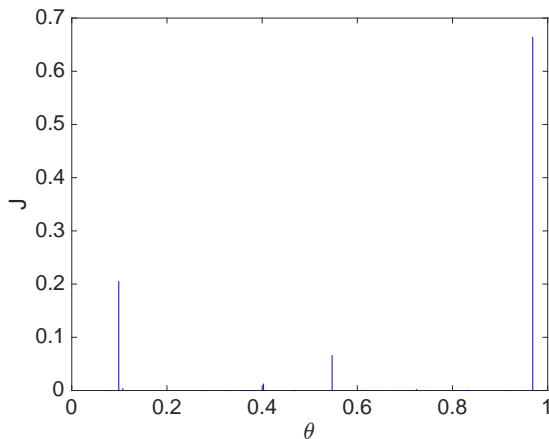


Gamma process simulation

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Gamma process simulation



In general, the Ferguson-Klass algorithm can be slow since $\eta(x)$ can only be numerically inverted for the gamma process.

Rejection method (Rosinski, 2001)

Suppose that we have two non-Gaussian Lévy process with Lévy intensities $\nu_1(x)$ and $\nu_2(x)$ with $\nu_1(x) < \nu_2(x)$ for all x .

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We can simulate a sequence z_1, z_2, \dots with intensity $\nu_1(x)$ by thinning a sequence y_1, y_2, \dots with intensity $\nu_2(x)$.

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Set $i = 1$ and $j = 1$

1. Simulate y_i from $\nu_2(x)$ using the Ferguson-Klass algorithm.

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Set $i = 1$ and $j = 1$

1. Simulate y_i from $\nu_2(x)$ using the Ferguson-Klass algorithm.
2. Simulate a uniform random variable u and calculate $\omega_i = \nu_1(y_i)/\nu_2(y_i)$. If $u < \omega_i$ set $z_j = y_i$, $j = j + 1$, $i = i + 1$. Otherwise, set $i = i + 1$.

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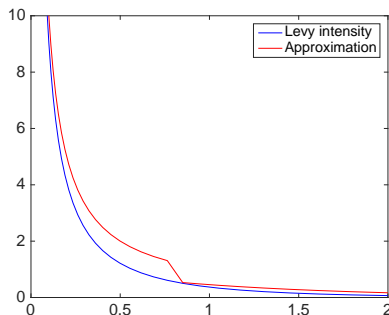
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3. Goto step 1.

Bounding functions for the gamma process (Griffin, 2019)

$$\nu_2(x) = \begin{cases} -\frac{1}{x} & x < b \\ b^{-1} \exp\{-x\} & x \geq b \end{cases},$$

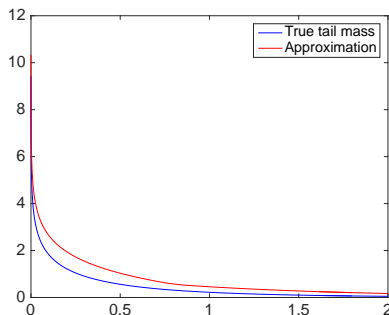
where $b = 0.8065$.



Bounding functions for the gamma process

$$\eta_2(x) = \begin{cases} -\log x + \log b + b^{-1} \exp\{-b\} & x < b \\ b^{-1} \exp\{-x\} & x \geq b \end{cases},$$

where $b = 0.8065$.



Truncation methods for gamma processes

1. Normalized gamma random variables

$$w_j = \frac{\gamma_j}{\sum_{k=1}^K \gamma_k} \text{ where } \gamma_k \sim \text{Ga}(M/K, 1).$$

Truncation methods for gamma processes

1. Normalized gamma random variables
 $w_j = \frac{\gamma_j}{\sum_{k=1}^K \gamma_k}$ where $\gamma_k \sim \text{Ga}(M/K, 1)$.
2. Ferguson and Klass representation
Choose a finite number of jumps K

$$F = \sum_{k=1}^K J_k \delta_{\theta_k}$$

Note: the jumps are ordered $J_1 > J_2 > J_3 > \dots$

3. ϵ -approximation

$$F = J_0 \delta_{\theta_0} + \sum_{k=1}^{K_\epsilon} J_k \delta_{\theta_k}$$

where $J_1, \dots, J_{K_\epsilon}$ are **all** jumps greater than ϵ .

$K_\epsilon \sim \text{Pn}(\int_\epsilon^\infty \nu(x) dx)$ (also, $K_\epsilon \sim \text{Pn}(\eta(\epsilon))$) and

$$p(J_k) = \frac{\nu(J_k)}{\int_\epsilon^\infty \nu(x) dx}, \quad J_k > \epsilon \text{ for } k = 0, 1, \dots, K_\epsilon.$$

Note: $\sum_{k=1}^{K_\epsilon} J_k \delta_{\theta_k}$ is a compound Poisson process and the truncation error (without J_0) is a Lévy process.

Illustration of ϵ -truncation

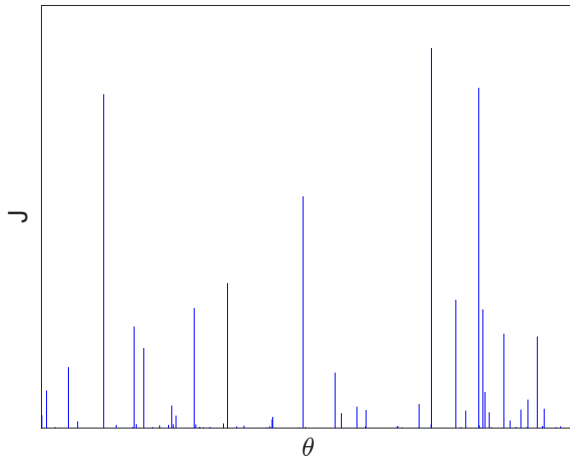


Illustration of ϵ -truncation

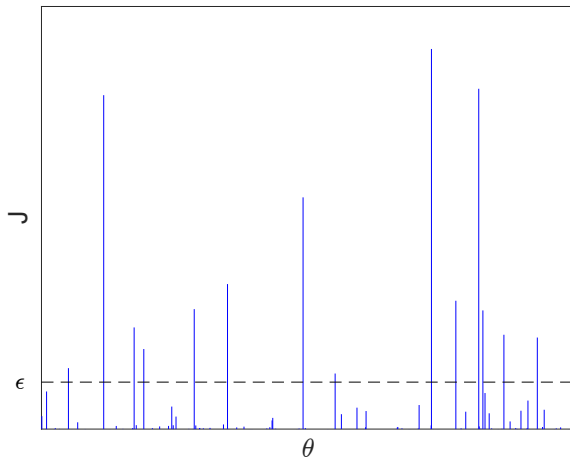
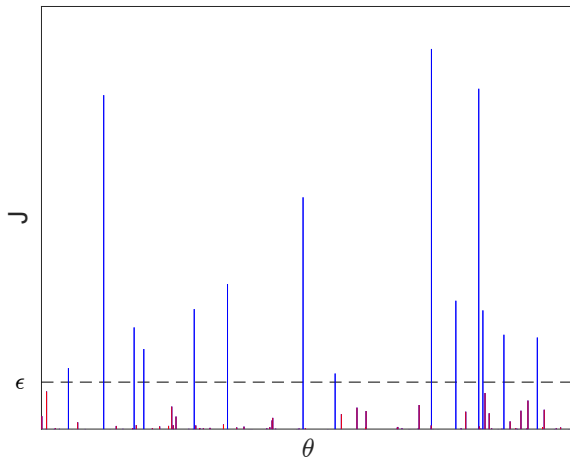


Illustration of ϵ -truncation



Sampling using normalized gamma random variables

The posterior distribution of $s_1, \dots, s_n, \gamma_1, \dots, \gamma_K, \theta_1, \dots, \theta_K$ is proportional to

$$\prod_{k=1}^K p(\gamma_k) \prod_{k=1}^K h(\theta_k) \prod_{i=1}^n w_{s_i} \prod_{i=1}^n k(y_i | \theta_{s_i})$$

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$$\begin{aligned} & \prod_{k=1}^K p(\gamma_k) \prod_{k=1}^K h(\theta_k) \prod_{i=1}^n w_{s_i} \prod_{i=1}^n k(y_i | \theta_{s_i}) \\ &= \prod_{k=1}^K p(\gamma_k) \prod_{k=1}^K h(\theta_k) \frac{\prod_{i=1}^n \gamma_{s_i}}{\left(\sum_{j=1}^K \gamma_j\right)^n} \prod_{i=1}^n k(y_i | \theta_{s_i}) \end{aligned}$$

Sampling using normalized gamma random variables

The “e to the minus v” trick

$$\frac{1}{\Gamma(n)} \int v^{n-1} \exp\{-va\} dx = \frac{1}{a^n}$$

Sampling using normalized gamma random variables

The “e to the minus v” trick

$$\frac{1}{\Gamma(n)} \int v^{n-1} \exp\{-va\} dx = \frac{1}{a^n}$$

We define a posterior augmented with v which is proportional to

$$\prod_{k=1}^K p(\gamma_k) \prod_{k=1}^K h(\theta_k) \prod_{i=1}^n \gamma_{s_i} v^{n-1} \exp \left\{ -v \sum_{j=1}^K \gamma_j \right\} \prod_{i=1}^n k(y_i | \theta_{s_i})$$

The Gibbs sampler

The full conditional of the Gibbs sampler are

1. The full conditional distribution of ν is $\text{Ga}(n, \sum_{j=1}^K \gamma_j)$.
2. The full conditional distribution of γ_k is $\text{Ga}(M/K + n_k, 1 + \nu)$.
3. The full conditional distribution of s_i is a discrete distribution with

$$p(s_i = k) \propto \gamma_k k(y_i | \theta_k)$$

4. The full conditional distribution of θ_k has density proportional to

$$h(\theta_k) \prod_{\{i|s_i=k\}} k(y_i | \theta_{s_i})$$

Sampling using ϵ -truncation

An augment posterior is constructed in a similar way to before

$$p(v, s_1, \dots, s_n, J_0, \dots, J_{K_\epsilon}, \theta_0, \dots, \theta_{K_\epsilon}) \\ \propto \prod_{k=0}^{K_\epsilon} p(J_k) \prod_{k=0}^{K_\epsilon} h(\theta_k) \prod_{i=1}^n J_{s_i} v^{n-1} \exp \left\{ -v \sum_{k=0}^{K_\epsilon} J_k \right\} \prod_{i=1}^n k(y_i | \theta_{s_i})$$

The Gibbs sampler

The full conditional of the Gibbs sampler are

1. The full conditional distribution of v is $\text{Ga}(n, \sum_{k=0}^{K_\epsilon} J_k)$.
2. The full conditional distribution of s_i is a discrete distribution with

$$p(s_i = k) \propto J_k p(y_i | \theta_k)$$

3. The full conditional distribution of θ_k has density proportional to

$$h(\theta_k) \prod_{\{i | s_i = k\}} k(y_i | \theta_{s_i})$$

4. The standard theory for NRM prior can be used to define the following scheme. Let J' be the set of jumps with observations allocated them the

$$J'_k \sim \text{Ga}(n_k, 1 + \nu), \quad J'_k > \epsilon$$

where n_k is the number of observation allocated to the k -th jump in J' .

Let \tilde{J} be the jumps without allocated observations then the \tilde{J} follow an ϵ -approximation with Lévy intensity

$$\exp\{-\nu J\} \nu(J), \quad J > \epsilon.$$

Series representations

We have looked at a few series representation but there are many.

Campbell et al. (2019) provide an overview of previous work. See also Nguyen et al. (2021) and Lee et al. (2023).

Slice sampling NRM mixtures (Griffin and Walker, 2011)

It's necessary to introduce latent variables v_1, v_2, \dots, v_n to fit these models using slice sampling and define

$$p(y, u, v|s) \propto v^{n-1} \prod_{i=1}^n I(u_i < J_{s_i}) \exp \left\{ -v \sum_{k=1}^{\infty} J_k \right\} k(y_i | \theta_{s_i}).$$

which involves an infinite number of jumps.

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which involves an infinite number of jumps.

This is resolved by integrating out all jumps smaller than L where $L = \min\{u_i\}$ and define $J_1, J_2, \dots, J_K > L$ then use

$$v^{n-1} \prod_{i=1}^n I(u_i < J_{s_i}) \exp \left\{ -v \sum_{k=1}^K J_k \right\} \mathbb{E} \left[\exp \left\{ -v \sum_{k=K+1}^{\infty} J_k \right\} \right] k(y_i | \theta_{s_i}).$$

Slice sampling NRM mixtures

The expectation is

$$\mathbb{E} \left[\exp \left\{ -v \sum_{k=K+1}^{\infty} J_k \right\} \right] = \exp \left\{ - \int_0^L (1 - \exp\{-vx\}) \nu(x) dx \right\}$$

due to definition of a Lévy intensity.

Gibbs sampling

The method can be made efficient by jointly updating u and J using the results of James et al. (2009) in the following steps.

1. Let J' be the set of jumps with observations allocated to them and let $n_j = \#\{i' | s_i = j\}$ then the full conditional of $J_j \in J'$ is proportional to

$$J_j^{n_j} \exp\{-\nu J_j\} \nu(J_j)$$

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2. Simulate u_1, u_2, \dots, u_n where $u_i \sim U(0, J_{s_i})$ and set $L = \min\{u_i\}$.

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2. Simulate u_1, u_2, \dots, u_n where $u_i \sim U(0, J_{s_i})$ and set $L = \min\{u_i\}$.
3. Simulate the jumps with no observations allocated to them from a Poisson process on (L, ∞) with intensity function $\exp\{-\nu x\} \nu(x)$.

Gibbs sampling for Dirichlet process mixtures

The method can be made efficient by jointly updating u and J using the results of James et al. (2009) in the following steps.

Gibbs sampling for Dirichlet process mixtures

The method can be made efficient by jointly updating u and J using the results of James et al. (2009) in the following steps.

1. Let J' be the set of jumps with observations allocated to them and let $n_j = \#\{i | s_i = j\}$ then the full conditional of $J_j \in J'$ is proportional to $J_j^{n_j-1} \exp\{-(1+\nu)J_j\}$, i.e. $\text{Ga}(n_j, 1+\nu)$.

Gibbs sampling for Dirichlet process mixtures

The method can be made efficient by jointly updating u and J using the results of James et al. (2009) in the following steps.

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2. Simulate u_1, u_2, \dots, u_n where $u_i \sim U(0, J_{s_i})$ and set $L = \min\{u_i\}$.
3. Simulate the jumps with no observations allocated to them from a Poisson process on (L, ∞) with intensity function $Mx^{-1} \exp\{-(1+\nu)x\}$.

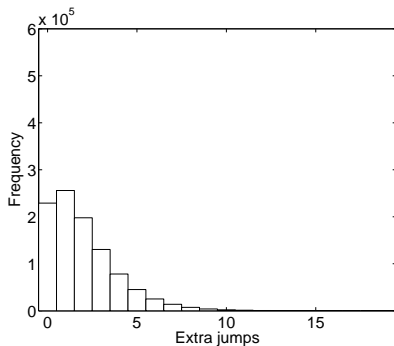
Slice 2

A slice sampler can also be defined with a single extra variable using the likelihood

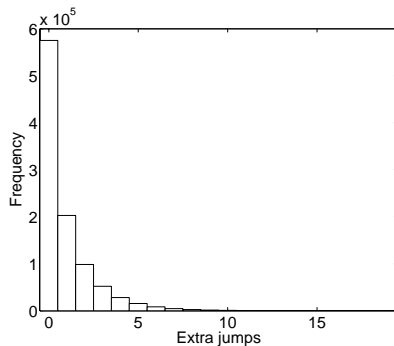
$$v^{n-1} \prod_{i=1}^n I(u < \min\{J_{s_i}\}) \frac{J_{s_i}}{\min\{J_{s_i}\}} \exp \left\{ -v \sum_{k=1}^{\infty} J_k \right\} k(y_i | \theta_{s_i}).$$

Results - Galaxy Data / MDP

Slice 1

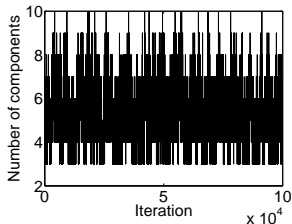


Slice 2

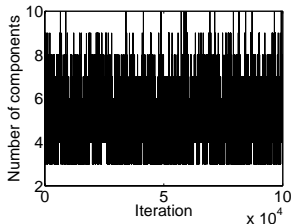


Comparisons - Galaxy / MDP

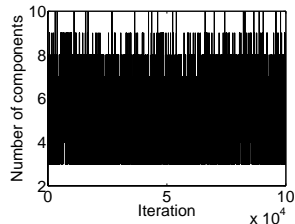
Slice 1



Slice 2



Auxiliary Gibbs Sampler



Results - Galaxy / NGG

a	Number of Clusters		Number of empty clusters	
	Mean	Standard Deviation	Mean	Standard Deviation
0.1	5.10	1.43	2.17	3.24
0.2	5.92	1.60	5.69	8.84
0.4	7.04	1.86	55.55	771.31

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