### Markov Chain Sampling Methods for Dirichlet Process Mixture Models

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#### Outline

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#### Introduction

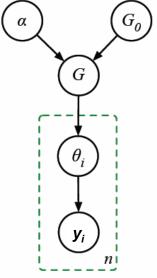


- Some problems are more accurately represented with non-conjugate priors
  - Audio interpolation (Godsill & Rayner, 1995)
  - Climatology opinion quantification (Al-Awadhi & Garthwaite, 2001)
  - Financial risk assessment (Siu & Yang, 1999)
- Non-conjugate priors + Gibbs = headache.
  - Update integrals are nasty to compute
- Solution? Metropolis-Hastings + partial Gibbs.



- Basic idea
  - Given data y<sub>1</sub>,...,y<sub>n</sub> ind. drawn from an unknown distribution (y<sub>i</sub> may be multivariate)
  - Model the unknown distribution as being drawn from of a mixture of distributions  $F(\theta)$ , w/ mixing distribution over  $\theta$  being G.
  - Let prior for G be a Dirichlet process w/ concentration parameter  $\alpha$  and base distribution  $G_0$ .
  - Then you have:

 $y_i \mid \theta_i \sim F(\theta_i)$  $\theta_i \mid G \sim G$  $G \sim D(G_0, \alpha)$ 





Integrate over G in previous model, giving a representation of the prior distribution of θ<sub>i</sub> in terms of previous θ's:

$$\theta_i \mid \theta_1, \dots, \theta_{i-1} \sim \frac{1}{i-1+\alpha} \sum_{j=1}^{i-1} \delta(\theta_j) + \frac{\alpha}{i-1+\alpha} G_0$$

- $\delta(\theta)$  is distribution concentrated at point  $\theta$ .
- You might notice the "Chinese Restaurant Process" at work here



You can also get here by letting K (# of components) go to ∞...

 $y_i \mid c, \phi \sim F(\phi_{c_i})$   $c_i \mid p \sim \text{Discrete}(p_1, \dots, p_K)$   $\phi_c \sim G_0$   $p_1, \dots, p_K \sim \text{Dirichlet}(\alpha/K, \dots, \alpha/K)$ 

- $c_i$  is the latent class associated with  $y_i$
- The parameters  $\varphi_c$  determine the distribution of observations from c



 Integrate over mixing proportions p<sub>c</sub> to write prior of c<sub>i</sub> as follows:

$$P(c_i = c \mid c_1, ..., c_{i-1}) = \frac{n_{i,c} + \alpha/K}{i - 1 + \alpha}$$

• Where  $n_{i,c}$  is the number of  $c_j$  for j < i equal to c. Letting K go to  $\infty$ , we get  $c_i$ 's prior as:  $P(c_i = c \mid c_1, \dots, c_{i-1}) \rightarrow \frac{n_{i,c}}{i-1+\alpha}$ 

$$P(c_i \neq c_j \text{ for all } j < i \mid c_1, \dots, c_{i-1}) \rightarrow \frac{\alpha}{i-1+\alpha}$$

# Gibbs sampling w/ conjugate priors

- Exact computation of posterior for DP mixture models not feasible, so use Monte Carlo approaches
- Sample from posterior of  $\theta_1, \ldots, \theta_n$  by simulating a Markov chain with this posterior as its equilibrium distribution
- Gibbs sampling is the natural approach here for conjugate priors
- 3 main ways of doing this

## Algorithm 1 (Escobar, 1994)



**Algorithm 1:** Let the state of the Markov chain consist of  $\theta_1, \ldots, \theta_n$ . Repeatedly sample as follows:

• For i = 1, ..., n: Draw a new value from  $\theta_i \mid \theta_{-i}, y_i$  as defined by equation (7).

$$\theta_i \mid \theta_{-i}, y_i \sim \sum_{j \neq i} q_{i,j} \,\delta(\theta_j) + r_i H_i$$

• Where  $H_i$  is the posterior for  $\theta$  based on the prior  $G_0$  and  $y_i$ , having likelihood  $F(y_i, \theta)$  and:

$$q_{i,j} = b F(y_i, \theta_j)$$
  
$$r_i = b \alpha \int F(y_i, \theta) dG_0(\theta)$$

• Convergence may be slow due to groups of observations that are highly probably to be associated with the same  $\theta$ 

# Algorithm 2 (West, Muller, & Escobar, 1994)



**Algorithm 2:** Let the state of the Markov chain consist of  $c_1, \ldots, c_n$  and  $\phi = (\phi_c : c \in \{c_1, \ldots, c_n\})$ . Repeatedly sample as follows:

- For i = 1, ..., n: If the present value of  $c_i$  is associated with no other observation (ie,  $n_{-i,c_i} = 0$ ), remove  $\phi_{c_i}$  from the state. Draw a new value for  $c_i$  from  $c_i | c_{-i}, y_i, \phi$  as defined by equation (11). If the new  $c_i$  is not associated with any other observation, draw a value for  $\phi_{c_i}$  from  $H_i$  and add it to the state.
- For all  $c \in \{c_1, \ldots, c_n\}$ : Draw a new value from  $\phi_c \mid y_i$  s.t.  $c_i = c$ .

If 
$$c = c_j$$
 for some  $j \neq i$ :  $P(c_i = c \mid c_{-i}, y_i, \phi) = b \frac{n_{-i,c}}{n-1+\alpha} F(y_i, \phi_c)$   
 $P(c_i \neq c_j \text{ for all } j \neq i \mid c_{-i}, y_i, \phi) = b \frac{\alpha}{n-1+\alpha} \int F(y_i, \phi) dG_0(\phi)$ 
(11)

### Algorithm 3 (Neal, 1992)



**Algorithm 3:** Let the state of the Markov chain consist of  $c_1, \ldots, c_n$ . Repeatedly sample as follows:

• For i = 1, ..., n: Draw a new value from  $c_i \mid c_{-i}, y_i$  as defined by equation (12).

If 
$$c = c_j$$
 for some  $j \neq i$ :  $P(c_i = c \mid c_{-i}, y_i) = b \frac{n_{-i,c}}{n-1+\alpha} \int F(y_i, \phi) dH_{-i,c}(\phi)$   
 $P(c_i \neq c_j \text{ for all } j \neq i \mid c_{-i}, y_i) = b \frac{\alpha}{n-1+\alpha} \int F(y_i, \phi) dG_0(\phi)$ 
(12)

### Methods for handling nonconjugate priors



- If *G*<sub>0</sub> is not the conjugate prior for *F*, the integrals for sampling from the posterior might not be feasible to compute.
- West, Muller, and Escobar suggested a Monte Carlo approximation to compute the integral (1994).
  - Slower convergence
  - New values of c<sub>i</sub> are likely to be discarded during following Gibbs iteration, leading to wrong distribution.

# Algorithm 4 (MacEachern & Muller, 1998)

**Algorithm 4:** Let the state of the Markov chain consist of  $c_1, \ldots, c_n$  and  $\phi = (\phi_c : c \in \{c_1, \ldots, c_n\})$ . Repeatedly sample as follows:

For i = 1,...,n: Let k<sup>-</sup> be the number of distinct c<sub>j</sub> for j≠i, and let these c<sub>j</sub> have values in {1,...,k<sup>-</sup>}. If c<sub>i</sub> ≠ c<sub>j</sub> for all j≠i, then with probability k<sup>-</sup>/(k<sup>-</sup>+1) do nothing, leaving c<sub>i</sub> unchanged. Otherwise, label c<sub>i</sub> as k<sup>-</sup>+1 if c<sub>i</sub> ≠ c<sub>j</sub> for all j≠i, or draw a value for φ<sub>k<sup>-</sup>+1</sub> from G<sub>0</sub> if c<sub>i</sub> = c<sub>j</sub> for some j≠i. Then draw a new value for c<sub>i</sub> from {1,...,k<sup>-</sup>+1} using the following probabilities:

$$P(c_i = c \mid c_{-i}, y_i, \phi_1, \dots, \phi_{k^-+1}) = \begin{cases} b n_{-i,c} F(y_i, \phi_c) & \text{if } 1 \le c \le k^- \\ b [\alpha/(k^-+1)] F(y_i, \phi_c) & \text{if } c = k^-+1 \end{cases}$$

where b is the appropriate normalizing constant. Change the state to contain only those  $\phi_c$  that are now associated with an observation.

• For all  $c \in \{c_1, \ldots, c_n\}$ : Draw a new value from  $\phi_c \mid y_i$  s.t.  $c_i = c$ , or perform some other update to  $\phi_c$  that leaves this distribution invariant.



### **Problem with Algorithm 4**



- Algorithm 4 has a problem in that assigning c<sub>i</sub> to a new component is reduced by a factor of k<sup>-</sup> + 1.
- However, something similar without this problem is possible.

# Metropolis-Hastings and partial Gibbs



- Use Metropolis-Hastings approach to update the c<sub>i</sub> using the conditional prior as the proposal distribution.
- Draw a candidate state, compute its acceptance probability. If it's accepted, use the candidate state, else leave as is.
- We can apply this to the finite model from slide 6, again integrating out *p<sub>c</sub>*

### Algorithm 5 (Neal, 1998)



**Algorithm 5:** Let the state of the Markov chain consist of  $c_1, \ldots, c_n$  and  $\phi = (\phi_c : c \in \{c_1, \ldots, c_n\})$ . Repeatedly sample as follows:

- For i = 1,...,n, repeat the following update of c<sub>i</sub> R times: Draw a candidate, c<sub>i</sub><sup>\*</sup>, from the conditional prior for c<sub>i</sub> given by equation (16). If a c<sub>i</sub><sup>\*</sup> not in {c<sub>1</sub>,..., c<sub>n</sub>} is proposed, chose a value for φ<sub>c<sub>i</sub><sup>\*</sup></sub> from G<sub>0</sub>. Compute the acceptance probability, a(c<sub>i</sub><sup>\*</sup>, c<sub>i</sub>), as in equation (15), and set the new value of c<sub>i</sub> to c<sub>i</sub><sup>\*</sup> with this probability. Otherwise let the new value of c<sub>i</sub> be the same as the old value.
- For all  $c \in \{c_1, \ldots, c_n\}$ : Draw a new value from  $\phi_c \mid y_i$  s.t.  $c_i = c$ , or perform some other update to  $\phi_c$  that leaves this distribution invariant.

$$a(c_i^*, c_i) = \min\left[1, \frac{F(y_i, \phi_{c_i^*})}{F(y_i, \phi_{c_i})}\right] \quad \text{If } c = c_j \text{ for some } j \neq i: P(c_i = c \mid c_{-i}) = \frac{n_{-i,c}}{n-1+\alpha}$$
$$P(c_i \neq c_j \text{ for all } j \neq i \mid c_{-i}) = \frac{\alpha}{n-1+\alpha}$$

## Algorithm 6 (Neal, 1998)



**Algorithm 6:** Let the state of the Markov chain consist of  $\theta_1, \ldots, \theta_n$ . Repeatedly sample as follows:

• For i = 1, ..., n, repeat the following update of  $\theta_i R$  times: Draw a candidate,  $\theta_i^*$ , from the following distribution:

$$\frac{1}{n-1+\alpha}\sum_{j\neq i}\,\delta(\theta_j) + \frac{\alpha}{n-1+\alpha}G_0$$

Compute the acceptance probability

$$a(\theta_i^*, \theta_i) = \min[1, F(y_i, \theta_i^*) / F(y_i, \theta_i)]$$

Set the new value of  $\theta_i$  to  $\theta_i^*$  with this probability; otherwise let the new value of  $\theta_i$  be the same as the old value.

#### Algorithm 7 (Neal, 1998)

**Algorithm 7:** Let the state of the Markov chain consist of  $c_1, \ldots, c_n$  and  $\phi = (\phi_c : c \in \{c_1, \ldots, c_n\})$ . Repeatedly sample as follows:

• For i = 1, ..., n, update  $c_i$  as follows: If  $c_i$  is a not a singleton (ie,  $c_i = c_j$  for some  $j \neq i$ ), let  $c_i^*$  be a newly-created component, with  $\phi_{c_i^*}$  drawn from  $G_0$ . Set the new  $c_i$  to this  $c_i^*$  with probability

$$a(c_i^*, c_i) = \min\left[1, \frac{\alpha}{n-1} \frac{F(y_i, \phi_{c_i^*})}{F(y_i, \phi_{c_i})}\right]$$

Otherwise, when  $c_i$  is a singleton, draw  $c_i^*$  from  $c_{-i}$ , choosing  $c_i^* = c$  with probability  $n_{-i,c} / (n-1)$ . Set the new  $c_i$  to this  $c_i^*$  with probability

$$a(c_i^*, c_i) = \min\left[1, \frac{n-1}{\alpha} \frac{F(y_i, \phi_{c_i^*})}{F(y_i, \phi_{c_i})}\right]$$

If the new  $c_i$  is not set to  $c_i^*$ , it is the same as the old  $c_i$ .

• For i = 1, ..., n: If  $c_i$  is a singleton (ie,  $c_i \neq c_j$  for all  $j \neq i$ ), do nothing. Otherwise, choose a new value for  $c_i$  from  $\{c_1, ..., c_n\}$  using the following probabilities:

$$P(c_i = c \mid c_{-i}, y_i, \phi, c_i \in \{c_1, \dots, c_n\}) = b \frac{n_{-i,c}}{n-1} F(y_i, \phi_c)$$

where b is the appropriate normalizing constant.

• For all  $c \in \{c_1, \ldots, c_n\}$ : Draw a new value from  $\phi_c \mid y_i$  s.t.  $c_i = c$ , or perform some other update to  $\phi_c$  that leaves this distribution invariant.



# Gibbs sampling w/ auxiliary parameters



- More flexible.
  - Basic idea is that we sample from a distribution  $\pi_x$  for x by sampling from distribution  $\pi_{xy}$  for (x, y).
  - Idea extendable to accommodate auxiliary variables which can be created/discarded during Markov chain simulation.
  - A variable y can be introduced temporarily:
    - Draw a value for y from its conditional given x
    - Perform an update of (x, y) leaving  $\pi_{xy}$  invariant
    - Discard y, leaving x.
  - This technique can be used to update c<sub>i</sub> for the DPM without having to integrate w.r.t. G<sub>0</sub>

#### Algorithm 8 (Neal, 1998)

**Algorithm 8:** Let the state of the Markov chain consist of  $c_1, \ldots, c_n$  and  $\phi = (\phi_c : c \in \{c_1, \ldots, c_n\})$ . Repeatedly sample as follows:

For i = 1,...,n: Let k<sup>-</sup> be the number of distinct c<sub>j</sub> for j≠i, and let h = k<sup>-</sup> + m. Label these c<sub>j</sub> with values in {1,...,k<sup>-</sup>}. If c<sub>i</sub> = c<sub>j</sub> for some j≠i, draw values independently from G<sub>0</sub> for those φ<sub>c</sub> for which k<sup>-</sup> < c ≤ h. If c<sub>i</sub> ≠ c<sub>j</sub> for all j≠i, let c<sub>i</sub> have the label k<sup>-</sup> + 1, and draw values independently from G<sub>0</sub> for those φ<sub>c</sub> for which k<sup>-</sup> + 1 < c ≤ h. Draw a new value for c<sub>i</sub> from {1,...,h} using the following probabilities:

$$P(c_{i} = c \mid c_{-i}, y_{i}, \phi_{1}, \dots, \phi_{h}) = \begin{cases} b \frac{n_{-i,c}}{n-1+\alpha} F(y_{i}, \phi_{c}) & \text{for } 1 \le c \le k^{-1} \\ b \frac{\alpha/m}{n-1+\alpha} F(y_{i}, \phi_{c}) & \text{for } k^{-1} < c \le h \end{cases}$$

where  $n_{-i,c}$  is the number of  $c_j$  for  $j \neq i$  that are equal to c, and b is the appropriate normalizing constant. Change the state to contain only those  $\phi_c$  that are now associated with one or more observations.

• For all  $c \in \{c_1, \ldots, c_n\}$ : Draw a new value from  $\phi_c \mid y_i$  s.t.  $c_i = c$ , or perform some other update to  $\phi_c$  that leaves this distribution invariant.



### **The Experiment**



	ne per iteration microseconds	Autocorrelation time for k	$\begin{array}{c} Autocorrelation \\ time \ for \ \theta_1 \end{array}$
Alg. 4 ("no gaps")	7.6	13.7	8.5
Alg. 5 (Metropolis-Hastings, $R = 4$ )	8.6	8.1	10.2
Alg. 6 (M-H, $R = 4$ , no $\phi$ update)	8.3	19.4	64.1
Alg. 7 (mod M-H & partial Gibbs)	8.0	6.9	5.3
Alg. 8 (auxiliary Gibbs, $m = 1$ )	7.9	5.2	5.6
Alg. 8 (auxiliary Gibbs, $m = 2$ )	8.8	3.7	4.7
Alg. 8 ( $m = 30$ , approximates Alg. 2	2) 38.0	2.0	2.8

- k is the number of distinct  $c_i$ ,  $\theta_1$  is the parameter associated with  $y_1$
- Algorithm 8 with m=1 superior to algorithm 4 ("no gaps")
- Performance much worse for algorithm 6, where no updates for  $\varphi_c$  are included
- With m=30, algorithm 8 takes longer, but performance is great.