

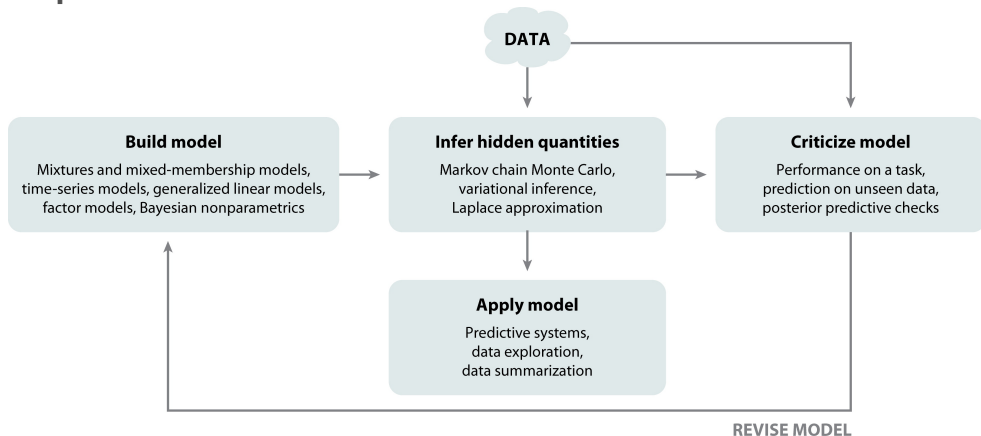
STATS271/371: Applied Bayesian Statistics

Bayesian Mixture Models and (Collapsed) Gibbs Sampling

Scott Linderman

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Box's Loop



Blei DM. 2014.

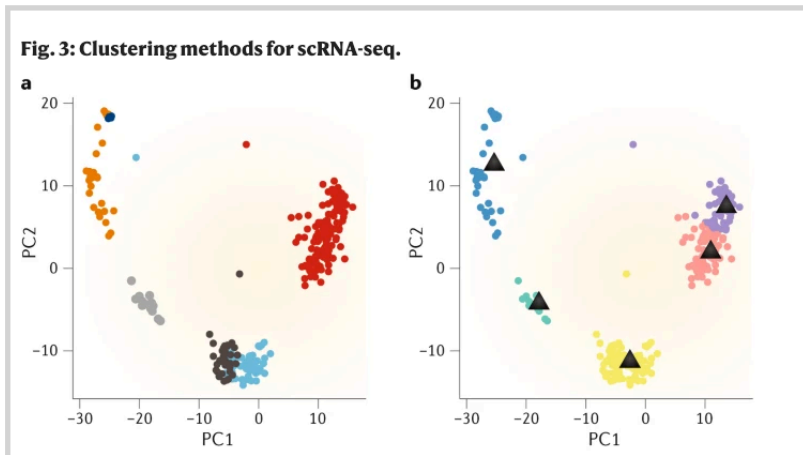
Annu. Rev. Stat. Appl. 1:203–32

Blei, *Ann. Rev. Stat. App.* 2014.

Lap 4: Bayesian Mixture Models and (Collapsed) Gibbs Sampling

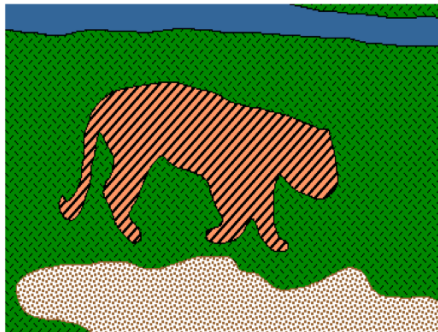
- ▶ **Model:** Bayesian mixture models
- ▶ **Algorithm:** Gibbs sampling
- ▶ **Criticism:** Posterior predictive checks
- ▶ **Algorithm II:** Collapsed Gibbs sampling

Motivation: Clustering scRNA-seq data



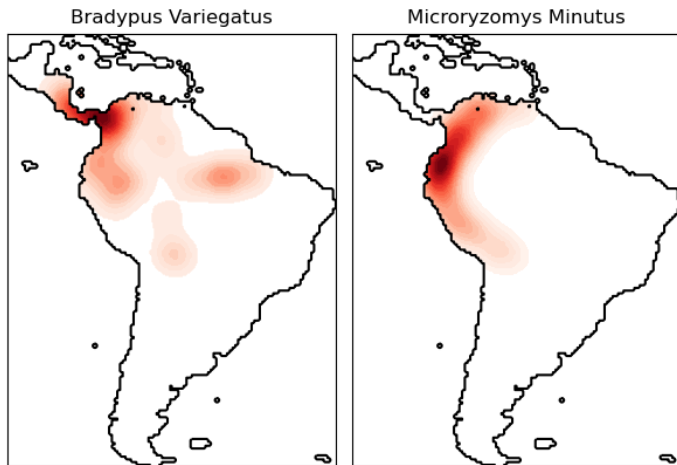
From Kiselev et al. [2019]

Motivation: Foreground/background segmentation



<https://ai.stanford.edu/~syyeung/cvweb/tutorial3.html>

Motivation: Density estimation



Notation

Constants: Let

- ▶ N denote the number of data points.
- ▶ K denote the number of mixture components (i.e. clusters)

Data: Let

- ▶ $\mathbf{x}_n \in \mathbb{R}^D$ denote the n -th data point.

Latent Variables: Let

- ▶ $z_n \in \{1, \dots, K\}$ denote the *assignment* of the n -th data point.

Notation II

Parameters: Let

- ▶ η_k denote the *natural parameters* of component k
- ▶ $\pi \in \Delta_K$ denote the component *proportions* (i.e. probabilities).

Hyperparameters: Let

- ▶ ϕ, ν denote hyperparameters of the prior on η
- ▶ $\alpha \in \mathbb{R}_+^K$ denote the concentration of the prior on proportions.

Generative Model

1. Sample the proportions from a Dirichlet prior:

$$\boldsymbol{\pi} \sim \text{Dir}(\boldsymbol{\alpha}) \quad (1)$$

2. Sample the parameters for each component:

$$\boldsymbol{\eta}_k \stackrel{\text{iid}}{\sim} p(\boldsymbol{\eta} \mid \boldsymbol{\phi}, \nu) \quad \text{for } k = 1, \dots, K \quad (2)$$

3. Sample the assignment of each data point:

$$z_n \stackrel{\text{iid}}{\sim} \boldsymbol{\pi} \quad \text{for } n = 1, \dots, N \quad (3)$$

4. Sample data points given their assignments:

$$\mathbf{x}_n \sim p(\mathbf{x} \mid \boldsymbol{\eta}_{z_n}) \quad \text{for } n = 1, \dots, N \quad (4)$$

Joint distribution

This generative model corresponds to the following factorization of the joint distribution,

$$p(\boldsymbol{\pi}, \{\boldsymbol{\eta}_k\}_{k=1}^K, \{(z_n, \mathbf{x}_n)\}_{n=1}^N \mid \boldsymbol{\phi}, \nu, \boldsymbol{\alpha}) = p(\boldsymbol{\pi} \mid \boldsymbol{\alpha}) \prod_{k=1}^K p(\boldsymbol{\eta}_k \mid \boldsymbol{\phi}, \nu) \prod_{n=1}^N p(z_n \mid \boldsymbol{\pi}) p(\mathbf{x}_n \mid z_n, \{\boldsymbol{\eta}_k\}_{k=1}^K) \quad (5)$$

Equivalently,

$$p(\boldsymbol{\pi}, \{\boldsymbol{\eta}_k\}_{k=1}^K, \{(z_n, \mathbf{x}_n)\}_{n=1}^N \mid \boldsymbol{\phi}, \nu, \boldsymbol{\alpha}) = p(\boldsymbol{\pi} \mid \boldsymbol{\alpha}) \prod_{k=1}^K p(\boldsymbol{\eta}_k \mid \boldsymbol{\phi}, \nu) \prod_{n=1}^N \prod_{k=1}^K [\Pr(z_n = k \mid \boldsymbol{\pi}) p(\mathbf{x}_n \mid \boldsymbol{\eta}_k)]^{\mathbb{I}[z_n=k]} \quad (6)$$

Substituting in the assumed forms

$$p(\boldsymbol{\pi}, \{\boldsymbol{\eta}_k\}_{k=1}^K, \{(z_n, \mathbf{x}_n)\}_{n=1}^N \mid \boldsymbol{\phi}, \boldsymbol{\alpha}) = \text{Dir}(\boldsymbol{\pi} \mid \boldsymbol{\alpha}) \prod_{k=1}^K p(\boldsymbol{\eta}_k \mid \boldsymbol{\phi}, \nu) \prod_{n=1}^N \prod_{k=1}^K [\pi_k p(\mathbf{x}_n \mid \boldsymbol{\eta}_k)]^{\mathbb{I}[z_n=k]} \quad (7)$$

Exponential family mixture models

What about $p(\mathbf{x} \mid \boldsymbol{\eta}_k)$ and $p(\boldsymbol{\eta}_k \mid \boldsymbol{\phi}, \nu)$?

Recall the *exponential family* distributions from Lap 2. Let's assume an exponential family likelihood,

$$p(\mathbf{x} \mid \boldsymbol{\eta}_k) = h(\mathbf{x}_n) \exp \left\{ \langle t(\mathbf{x}_n), \boldsymbol{\eta}_k \rangle - A(\boldsymbol{\eta}_k) \right\}. \quad (8)$$

Then assume a *conjugate prior*,

$$p(\boldsymbol{\eta}_k \mid \boldsymbol{\phi}, \nu) \propto \exp \left\{ \langle \boldsymbol{\phi}, \boldsymbol{\eta}_k \rangle - \nu A(\boldsymbol{\eta}_k) \right\}. \quad (9)$$

The hyperparameters $\boldsymbol{\phi}$ are *pseudo-observations* of the sufficient statistics (like statistics from fake data points) and ν is a *pseudo-count* (like the number of fake data points).

Note that the product of prior and likelihood remains in the same family as the prior. That's why we call it conjugate.

Example: Gaussian mixture model

Assume the conditional distribution of \mathbf{x}_n is a Gaussian with mean $\boldsymbol{\eta}_{z_n} \in \mathbb{R}^D$ and identity covariance,

$$p(\mathbf{x}_n | \boldsymbol{\eta}_k) = \mathcal{N}(\mathbf{x}_n | \boldsymbol{\eta}_k, I) \quad (10)$$

$$= (2\pi)^{-D/2} \exp \left\{ -\frac{1}{2} (\mathbf{x}_n - \boldsymbol{\eta}_k)^\top (\mathbf{x}_n - \boldsymbol{\eta}_k) \right\} \quad (11)$$

$$= (2\pi)^{-D/2} \exp \left\{ -\frac{1}{2} \mathbf{x}_n^\top \mathbf{x}_n + \mathbf{x}_n^\top \boldsymbol{\eta}_k - \frac{1}{2} \boldsymbol{\eta}_k^\top \boldsymbol{\eta}_k \right\}, \quad (12)$$

which is an exponential family distribution with base measure $h(\mathbf{x}_n) = (2\pi)^{-D/2} e^{-\frac{1}{2} \mathbf{x}_n^\top \mathbf{x}_n}$, sufficient statistics $t(\mathbf{x}_n) = \mathbf{x}_n$, and log normalizer $A(\boldsymbol{\eta}_k) = \frac{1}{2} \boldsymbol{\eta}_k^\top \boldsymbol{\eta}_k$.

Then assume a Gaussian prior on the component parameters. It's conjugate,

$$p(\boldsymbol{\eta}_k | \boldsymbol{\phi}, \nu) = \mathcal{N}(\nu^{-1} \boldsymbol{\phi}, \nu^{-1} I) \propto \exp \left\{ \boldsymbol{\phi}^\top \boldsymbol{\eta}_k - \frac{\nu}{2} \boldsymbol{\eta}_k^\top \boldsymbol{\eta}_k \right\} = \exp \left\{ \boldsymbol{\phi}^\top \boldsymbol{\eta}_k - \nu A(\boldsymbol{\eta}_k) \right\}. \quad (13)$$

Note that $\boldsymbol{\phi}$ sets the location and ν sets the precision (i.e. inverse variance).

MAP inference via coordinate ascent

Before diving into fully Bayesian inference algorithms, let's first consider **MAP inference**.

Idea: find the mode of $p(\pi, \{\eta_k\}_{k=1}^K, \{z_n\}_{n=1}^N \mid \{\mathbf{x}_n\}_{n=1}^N, \phi, \nu, \alpha)$ by **coordinate ascent**.

For now, set $\phi = \mathbf{0}$, and $\nu = 0$ so that the prior is an (improper) uniform distribution. Then maximizing the posterior is equivalent to maximizing the likelihood.

While we're simplifying, let's even fix $\pi = \frac{1}{K} \mathbf{1}_K$.

Coordinate ascent in the Gaussian mixture model

For the Gaussian mixture model (with uniform prior and $\pi = \frac{1}{K}\mathbf{1}_K$), coordinate ascent amounts to:

1. For each $n = 1, \dots, N$, fix all variables but z_n and find z_n^\star that maximizes

$$p(\pi, \{\boldsymbol{\eta}_k\}_{k=1}^K, \{(z_n, \mathbf{x}_n)\}_{n=1}^N \mid \boldsymbol{\phi}, \nu, \boldsymbol{\alpha}) \propto p(\mathbf{x}_n \mid z_n, \{\boldsymbol{\eta}_k\}_{k=1}^K) = \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\eta}_{z_n}, I) \quad (14)$$

The cluster assignment that maximizes the likelihood is the one with the closest mean to \mathbf{x}_n , so set

$$z_n^\star = \arg \min_{k \in \{1, \dots, K\}} \|\mathbf{x}_n - \boldsymbol{\eta}_k\|_2. \quad (15)$$

Coordinate ascent in the Gaussian mixture model II

2 For each $k = 1, \dots, K$, fix all variables but $\boldsymbol{\eta}_k$ and find $\boldsymbol{\eta}_k^*$ that maximizes,

$$p(\boldsymbol{\pi}, \{\boldsymbol{\eta}_k\}_{k=1}^K, \{(z_n, \mathbf{x}_n)\}_{n=1}^N \mid \boldsymbol{\phi}, \nu, \boldsymbol{\alpha}) \propto \prod_{n=1}^N p(\mathbf{x}_n \mid \boldsymbol{\eta}_k)^{\mathbb{I}[z_n=k]} \quad (16)$$

$$\propto \exp \left\{ \sum_{n=1}^N \mathbb{I}[z_n = k] \left(\mathbf{x}_n^\top \boldsymbol{\eta}_k - \frac{1}{2} \boldsymbol{\eta}_k^\top \boldsymbol{\eta}_k \right) \right\} \quad (17)$$

Taking the derivative of the log and setting to zero yields,

$$\boldsymbol{\eta}_k^* = \frac{1}{N_k} \sum_{n=1}^K \mathbb{I}[z_n = k] \mathbf{x}_n, \quad (18)$$

where $N_k = \sum_{n=1}^N \mathbb{I}[z_n = k]$.

This is the **k-means algorithm**!

Aside: EM in the Gaussian mixture model

We'll talk more about *coordinate ascent variational inference* (CAVI) and *expectation-maximization* (EM) next week. Not to spoil the surprise, but we'll see that they have a similar flavor. Instead of assigning z_n^* to the closest cluster, we compute *responsibilities* for each cluster:

1. For each data point n and component k , set the *responsibility* to,

$$\omega_{nk} = \frac{\pi_k \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\eta}_k, I)}{\sum_{j=1}^K \pi_j \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\eta}_j, I)}. \quad (19)$$

2. For each component k , set the mean to

$$\boldsymbol{\eta}_k^* = \frac{1}{N_k} \sum_{n=1}^K \omega_{nk} \mathbf{x}_n, \quad (20)$$

where $N_k = \sum_{n=1}^N \omega_{nk}$.

Note that EM allows for arbitrary proportions π . Those can be updated as well: for each component k , set $\pi_k = \frac{N_k}{N}$.

Lap 4: Bayesian Mixture Models and (Collapsed) Gibbs Sampling

- ▶ Model: Bayesian mixture models
- ▶ **Algorithm: Gibbs sampling**
- ▶ Criticism: Posterior predictive checks
- ▶ Algorithm II: Collapsed Gibbs sampling

Gibbs sampling in Bayesian mixture models

Idea: just like in coordinate ascent, update one variable at a time. *But rather than setting it to its conditional mode, sample from its conditional distribution.*

Gibbs sampling in Bayesian mixture models II

1. For each data point n , sample a new assignment from the complete conditional distribution

$$z_n \sim p(z_n \mid \{\mathbf{x}_n\}_{n=1}^N, \{z_{n'}\}_{n' \neq n}, \{\boldsymbol{\eta}_k\}_{k=1}^K, \boldsymbol{\pi}, \boldsymbol{\phi}, \nu, \boldsymbol{\alpha}). \quad (21)$$

Thanks to the factorization of the joint distribution,

$$\Pr(z_n = k \mid -) \propto \Pr(z_n = k \mid \boldsymbol{\pi}) p(x_n \mid \boldsymbol{\eta}_k) \quad (22)$$

In the Gaussian mixture model, this is,

$$\Pr(z_n = k \mid -) = \frac{\pi_k \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\eta}_k, I)}{\sum_{j=1}^K \pi_j \mathcal{N}(\mathbf{x}_n \mid \boldsymbol{\eta}_j, I)} \quad (23)$$

$$\equiv \omega_{nk}. \quad (24)$$

I.e., Gibbs sampling generates *random* assignments by sampling according to the responsibilities.

Gibbs sampling in Bayesian mixture models III

2 For each component k , sample new parameters from their complete conditional,

$$\boldsymbol{\eta}_k \sim p(\boldsymbol{\eta}_k \mid \{(\mathbf{x}_n, z_n)\}_{n=1}^N, \{\boldsymbol{\eta}_{k'}\}_{k' \neq k}, \boldsymbol{\pi}, \boldsymbol{\phi}, \nu, \boldsymbol{\alpha}). \quad (25)$$

Thanks to the factorization of the joint distribution,

$$p(\boldsymbol{\eta}_k \mid -) \propto p(\boldsymbol{\eta}_k \mid \boldsymbol{\phi}, \nu) \prod_{n: z_n=k} p(x_n \mid \boldsymbol{\eta}_k). \quad (26)$$

In an Gaussian mixture model,

$$p(\boldsymbol{\eta}_k \mid -) \propto \exp \left\{ \left(\boldsymbol{\phi} + \sum_{n=1}^N \mathbb{I}[z_n = k] \mathbf{x}_n \right)^\top \boldsymbol{\eta}_k - \frac{\nu + N_k}{2} \boldsymbol{\eta}_k^\top \boldsymbol{\eta}_k \right\} \quad (27)$$

$$\propto \mathcal{N} \left(\boldsymbol{\eta}_k \mid (\nu + N_k)^{-1} \left(\boldsymbol{\phi} + \sum_{n=1}^N \mathbb{I}[z_n = k] \mathbf{x}_n \right), (\nu + N_k)^{-1} \mathbf{I} \right) \quad (28)$$

where $N_k = \sum_{n=1}^N \mathbb{I}[z_n = k]$. What happens when $N_k \rightarrow \infty$?

Gibbs sampling in Bayesian mixture models IV

3 Finally, sample new component proportions from their complete conditional,

$$\pi \sim p(\pi \mid \{(\mathbf{x}_n, z_n)\}_{n=1}^N, \{\boldsymbol{\eta}_k\}_{k=1}^K, \boldsymbol{\phi}, \nu, \boldsymbol{\alpha}). \quad (29)$$

Thanks to the factorization of the joint distribution,

$$p(\pi \mid -) \propto \text{Dir}(\pi \mid \boldsymbol{\alpha}) \prod_{n=1}^N p(z_n \mid \pi) \quad (30)$$

$$\propto \prod_{k=1}^K \pi_k^{\alpha_k - 1} \times \prod_{n=1}^N \prod_{k=1}^K \pi_k^{\mathbb{I}[z_n = k]} \quad (31)$$

$$\propto \text{Dir}(\pi \mid [\alpha_1 + N_1, \dots, \alpha_K + N_K]) \quad (32)$$

where $N_k = \sum_{n=1}^N \mathbb{I}[z_n = k]$. What happens when $N_k \rightarrow \infty$?

Gibbs sampling in Bayesian exponential family mixture models

What happens in general exponential family models? Step 2 becomes,

2 For each component k , sample new parameters from their complete conditional,

$$p(\boldsymbol{\eta}_k | -) \propto \exp \left\{ \left\langle \boldsymbol{\phi} + \sum_{n=1}^N \mathbb{I}[z_n = k] t(\mathbf{x}_n), \boldsymbol{\eta}_k \right\rangle - (\nu + N_k) A(\boldsymbol{\eta}_k) \right\} \quad (33)$$

$$= p \left(\boldsymbol{\eta}_k \mid \boldsymbol{\phi} + \sum_{n=1}^N \mathbb{I}[z_n = k] t(\mathbf{x}_n), \nu + N_k \right) \quad (34)$$

where $N_k = \sum_{n=1}^N \mathbb{I}[z_n = k]$.

Opportunities for parallelism

- ▶ In all three algorithms above, the updates of z_n are independent of one another (once you fix $\{\eta_k\}_{k=1}^K$) and hence can be performed in parallel.
- ▶ Likewise, the updates of η_k are independent of one another (once you fix $\{z_n\}_{n=1}^N$) and hence can be performed in parallel.
- ▶ In fact, we can write these as simple map-reduce algorithms and take advantage of parallel hardware if it's available.
- ▶ In the Gibbs sampling case, updating many variables at once from their combined conditional distribution is called **blocked Gibbs sampling**, and it's particularly easy when the variables are conditionally independent, as in the Bayesian mixture model.

Next time

- ▶ Show that **Gibbs is a special case of MH** and, as such, asymptotically generates samples from the posterior distribution.
- ▶ Talk about **posterior predictive checks** and ways of choosing K .
- ▶ Introduce **collapsed Gibbs sampling**, which will allow us to generalize to **nonparametric Bayesian mixture models**.

References I

Vladimir Yu Kiselev, Tallulah S Andrews, and Martin Hemberg. Challenges in unsupervised clustering of single-cell RNA-seq data. *Nat. Rev. Genet.*, 20(5):273–282, May 2019.