



BROWN

# Memoized Online Variational Inference for Dirichlet Process Mixture Models

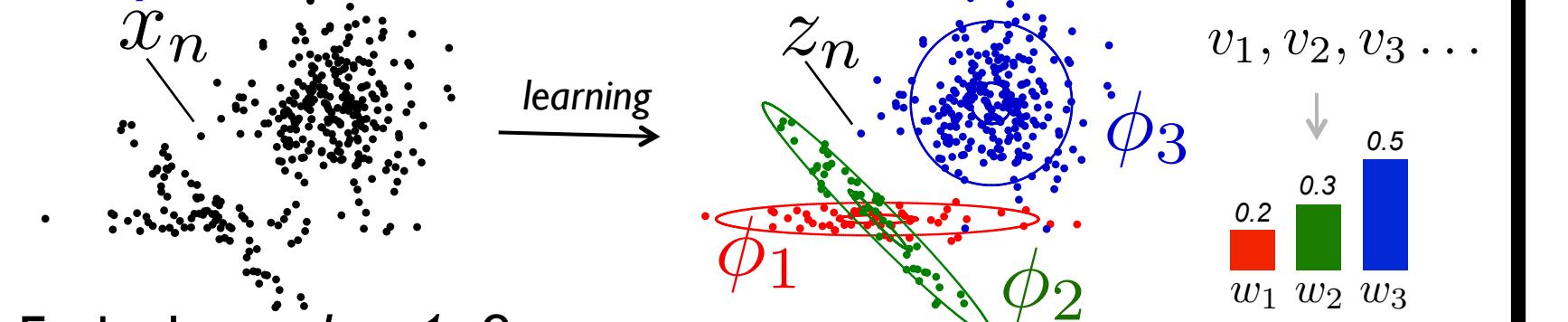
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Python code: [bitbucket.org/michaelchughes/bnpy/](https://bitbucket.org/michaelchughes/bnpy/)

## Dirichlet Process Mixture Model

Assigns data to discrete clusters

**Nonparametric:** number of clusters learned from data.



Each cluster  $k = 1, 2, \dots$ :

Stick fraction

$$v_k \sim \text{Beta}(1, \alpha_0)$$

Appearance probability

$$w_k = v_k \prod_{\ell=1}^{k-1} (1 - v_\ell)$$

stick-breaking

Data-generating parameter

$$\phi_k \sim H(\lambda_0)$$

Each data item  $n = 1, 2, \dots N$ :

Draw cluster assignment

$$z_n \sim \text{Discrete}(w_1, w_2, \dots)$$

Draw observed data

$$x_n \sim F(\phi_{z_n}) = \exp(\phi_{z_n}^T t(x_n) - a(\phi_{z_n}))$$

exponential family

Algorithms generalize

to any likelihood  $F$ ,

not just Gaussian

**Multivariate Gaussian** likelihood  $F$

$$x_n \sim \mathcal{N}(\mu_{z_n}, \Lambda_{z_n}^{-1})$$

mean, precision matrix

sufficient statistics

## Summary

Memoized online (MO) variational inference

- No pesky learning rates, insensitive to batch size

New online moves add/remove clusters on-the-fly

- Birth:** add useful clusters, escape local optima

- Merge:** remove redundancy, improve speed

**MO-BM** (MO with births and merges):  
Scalable, robust exploration of nonparametric posterior.  
Start with just  $K=1$  cluster, grow as needed!

## Stochastic Online (SO)

At batch  $b$ , perform usual E-step, then

**Update** global factors via noisy gradient

$$\lambda_k^b \leftarrow \lambda_0 + \frac{N}{|\mathcal{B}_b|} s_k^b$$

**M-step** amplifies current batch

$$\lambda_k \leftarrow \rho_t \lambda_k^b + (1-\rho_t) \lambda_k$$

Gradient step natural gradients make updates simple

Many options ( $a, b, c, \dots$ ) for decay schedule.

Learning rate  $\rho_t$

$$\rho_t \leftarrow (d+t)^{-\kappa}$$

**Find** (local) optima of full-data objective in expectation.

Sensitive to learning rate schedule and batch size. Careful tuning required.

## Birth Moves

Escape poor solutions by adding useful clusters.

- Each move adds many clusters via fresh analysis of one cluster's data.

**Collect** targeted subsample

Original data

Subsample explained by 1

Create new clusters

Brand-new DP-GMM learned via Full VB on  $x'$

expected size of each cluster

1 2 3 4 5 6 7

Batch 1

Batch b

Batch b+1

Batch B

current position

0 0 0 0 0

0 0 0 0 0

0 0 0 0 0

batches not-yet updated do not use new clusters

Before (K=2)

After (K=7)

Original clusters remain unaltered.

Expansion possible via nested truncation.

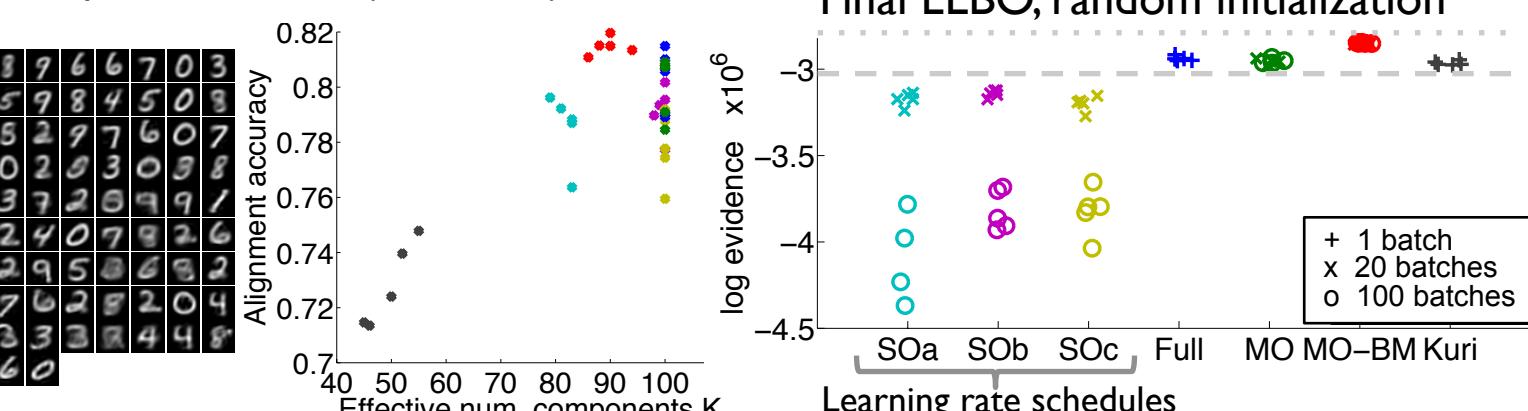
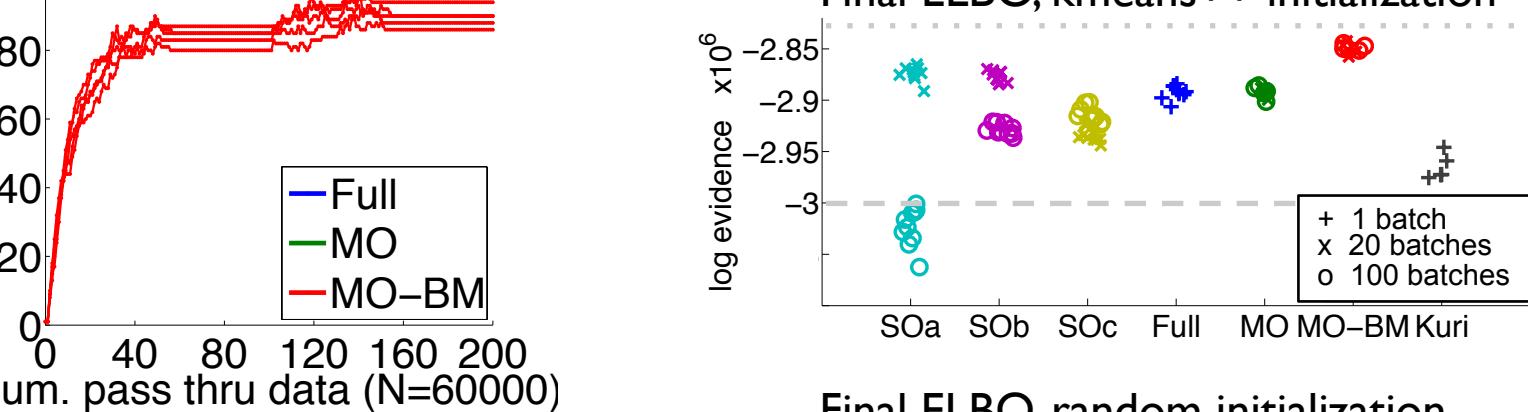
## MNIST Handwritten Digits

Cluster 60000 images of digits 0-9. PCA projected to 50 dimensions.

10 runs of each algorithm, from 10 fixed sets of initial parameters.

MO-BM started at  $K=1$  discovers >80 useful clusters via births

Final ELBO, kmeans++ initialization



MO-BM estimated clusters have best many-to-one alignment to true digits 0-9

MO reliable, while SO very sensitive to learning rate, # batches, & initialization

## Variational Bayes Inference (VB)

Algorithm that finds approximate posterior  $q$

- Coordinate ascent optimization, minimizes KL divergence

- Like EM, but learns distributions not just point estimates

**Truncation** to  $K$  clusters  $q(z_n > K) = 0$  is nested: allows  $K$  to grow/shrink

**Update** at each iteration

For data item  $n = 1, 2, \dots N$ :  
 $r_{n1} \dots r_{nK} \leftarrow \text{Estep}(x_n, \alpha, \lambda)$

$r_{nk} \propto e^{\log p(z_n=k|v) + \log p(x_n|z_n=k, \phi)}$

For cluster  $k = 1, 2, \dots K$ :

$s_k^0 \leftarrow \sum_{n=1}^N r_{nk} t(x_n)$  Expected sufficient stats  
 $\lambda_k \leftarrow \lambda_0 + s_k^0$  M-step

Process entire dataset between global updates.

Slow to propagate information.

Updates just simple function of  $\{N_k^0\}_{k=1}^K$

Evidence lower bound (ELBO) objective

$$\log p(x) \geq \mathcal{L}(q)$$

$$\mathcal{L}(q) = \sum_{k=1}^K \mathbb{E}[\phi_k]^T s_k^0 - N_k^0 \mathbb{E}[a(\phi_k)] + N_k^0 \mathbb{E}[\log r_{nk}] + \mathcal{L}(q(v), q(\phi))$$

linear function of sufficient statistics

**q(z) entropy** global factors

## Memoized Online (MO)

New variational algorithm, inspired by [Neal & Hinton '99]

- Online proposal, requires no batch processing

- Modest memory required, but still scales to millions of examples

- Several passes through all batches yield quality solutions

**Update** for each batch  $b$

$r(\mathcal{B}_b) \leftarrow \text{Estep}(x(\mathcal{B}_b), \alpha, \lambda)$

For cluster  $k = 1, 2, \dots K$ :

$s_k^0 \leftarrow s_k^0 - s_k^b$

$s_k^b \leftarrow \sum_{n \in \mathcal{B}_b} r_{nk} t(x_n)$  Expected sufficient stats

$s_k^0 \leftarrow s_k^0 + s_k^b$

$\lambda_k \leftarrow \lambda_0 + s_k^0$  M-step

Global factors updated at every batch.

$s_k^0 = s_k^1 + s_k^2 + \dots + s_k^B$

Global summaries are additive

**Data**

$x(\mathcal{B}_1)$

$x(\mathcal{B}_2)$

$\vdots$

$x(\mathcal{B}_b)$

$\vdots$

$x(\mathcal{B}_B)$

**Batch Summaries**

$s_1^1 s_2^1 \dots s_K^1$

$s_1^2 s_2^2 \dots s_K^2$

$\vdots$

$s_1^B s_2^B \dots s_K^B$

**Global Summary**

$s_1^0 s_2^0 \dots s_K^0$

## Merge Moves

Merge two clusters into one. Simpler models & faster learning.

- Run many proposals after each pass.

Online proposal, requires no batch processing

Modest memory required, but still scales to millions of examples

Several passes through all batches yield quality solutions

Accept/reject decision via exact, full-dataset ELBO comparison

accept if  $\mathcal{L}(q_{merge}) > \mathcal{L}(q)$

Requires cached entropy  $H_{k_a, k_b}^b$  for all pairs.

New cluster takes over all responsibility for any data assigned to old clusters.

$$r_{nk_m} \leftarrow r_{nk_a} + r_{nk_b}$$

Direct construction of global summaries:

$$s_{k_m}^0 \leftarrow s_{k_a}^0 + s_{k_b}^0$$

additivity

Accept/reject decision via exact, full-dataset ELBO comparison

accept if  $\mathcal{L}(q_{merge}) > \mathcal{L}(q)$

Requires cached entropy  $H_{k_a, k_b}^b$  for all pairs.

5x5 image patches, with strong edges

K=8 true clusters

$\mathcal{L}(q)$

MO K=25 Full K=25

MO-BM K=1

Accept/reject merge via exact full-dataset ELBO.

GreedyMerge: Only use current batch ELBO to accept/reject.

Global summaries are additive

All MO-BM runs find ideal, others local optima.

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