

# Mixture models for cluster analysis: from model-based inference to Bayesian non-parametrics

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# Outline

- 1 Model-based clustering
- 2 Parsimonious Gaussian mixture models
- 3 The Bayesian mixture for model-based clustering
- 4 The Bayesian non-parametric GMM (Infinite GMM)
- 5 Infinite parsimonious GMMs and Dirichlet Process Mixture
- 6 experiments
- 7 Conclusion

# Model-Based Clustering

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# EM algorithm

- where

$$\tau_{ik}^{(q)} = p(z_i = k | \mathbf{x}_i; \Psi^{(q)}) = \frac{\pi_k f_k(\mathbf{x}_i; \Psi_k^{(q)})}{\sum_{\ell=1}^K \pi_\ell f_\ell(\mathbf{x}_i; \Psi_\ell^{(q)})}$$

is the posterior probability that  $\mathbf{x}_i$  originates from the  $k$ th component density.

- In  $\mathbb{E}[z_{ik} | \mathbf{x}_i, \Psi^{(q)}]$ , we used the fact that conditional expectations and conditional probabilities are the same for the indicator binary-valued variables  $z_{ik}$ :  $\mathbb{E}[z_{ik} | \mathbf{x}_i, \Psi^{(q)}] = p(z_{ik} = 1 | \mathbf{x}_i, \Psi^{(q)})$ .

⇒ From the expression of  $Q(\Psi, \Psi^{(q)})$ , we can see that this step simply requires the computation of the posterior probabilities  $\tau_{ik}^{(q)}$ .













# EM for GMMs

- The observed-data log-likelihood of  $\Psi$  for the Gaussian mixture model:

$$\mathcal{L}(\Psi; \mathbf{X}) = \sum_{i=1}^n \log \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

- The complete-data log-likelihood of  $\Psi$  for the Gaussian mixture model:

$$\mathcal{L}_c(\Psi; \mathbf{X}, \mathbf{z}) = \sum_{i=1}^n \sum_{k=1}^K z_{ik} \log \pi_k \mathcal{N}(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

EM:

- Starts with an initial parameter  $\Psi^{(0)} = (\pi_1^{(0)}, \dots, \pi_K^{(0)}, \Psi_1^{(0)}, \dots, \Psi_K^{(0)})$  where  $\Psi_k^{(0)} = (\boldsymbol{\mu}_k^{(0)}, \boldsymbol{\Sigma}_k^{(0)})$

# E-Step for GMMs

- the expected complete-data log-likelihood:

$$\begin{aligned} Q(\Psi, \Psi^{(q)}) &= \mathbb{E} \left[ \mathcal{L}_c(\Psi; \mathbf{X}, \mathbf{z}) | \mathbf{X}; \Psi^{(q)} \right] \\ &= \sum_{i=1}^n \sum_{k=1}^K \tau_{ik}^{(q)} \log \pi_k + \sum_{i=1}^n \sum_{k=1}^K \tau_{ik}^{(q)} \log \mathcal{N}(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k). \end{aligned}$$

⇒ This step therefore computes the posterior probabilities

$$\tau_{ik}^{(q)} = p(z_i = k | \mathbf{x}_i, \Psi^{(q)}) = \frac{\pi_k \mathcal{N}(\mathbf{x}_i; \boldsymbol{\mu}_k^{(q)}, \boldsymbol{\Sigma}_k^{(q)})}{\sum_{\ell=1}^K \pi_\ell \mathcal{N}(\mathbf{x}_i; \boldsymbol{\mu}_\ell^{(q)}, \boldsymbol{\Sigma}_\ell^{(q)})}$$

that  $\mathbf{x}_i$  originates from the  $k$ th component density.





# Initialization Strategies and stopping rules for EM

- The initialization of EM is a crucial point since it maximizes locally the log-likelihood.
- if the initial value is inappropriately selected, the EM algorithm may lead to an unsatisfactory estimation.
- The most used strategy: use several EM tries and select the solution maximizing the log-likelihood among those runs.
- For each run of EM, one can initialize it
  - randomly
  - by Computing a parameter estimate from another clustering algorithm such as  $K$ -means, Classification EM, Stochastic EM ...
  - with a few number of steps of EM itself.
- Stop EM when the relative increase of the log-likelihood between two iterations is below a fixed threshold  $|\frac{\mathcal{L}^{(q+1)} - \mathcal{L}^{(q)}}{\mathcal{L}^{(q)}}| \leq \epsilon$  or when a predefined number of iterations is reached.

# EM properties

- The EM algorithm always monotonically increases the observed-data log-likelihood.
- The sequence of parameter estimates generated by the EM algorithm converges toward at least a local maximum or a stationary value of the incomplete-data likelihood function.
- numerical stability
- simplicity of implementation
- reliable convergence
- In general, both the E- and M-steps will have particularly simple forms when the complete-data probability density function is from the exponential family;
- Some drawbacks: EM is sometimes very slow to converge especially for high dimensional data;  
in some problems, the E- or M-step may be analytically intractable (but this can be tackled by using EM extensions)

# EM extensions

- The EM variants mainly aim at:
  - ① increasing the convergence speed of EM and addressing the optimization problem in the M-step
  - ② computing the E-step when it is intractable.
- In the first case, one can speak about deterministic algorithms :
  - e.g., Incremental EM (IEM)
  - Gradient EM
  - Generalized EM (GEM) algorithm
  - Expectation Conditional Maximization (ECM)
  - Expectation Conditional Maximization Either (ECME)
- In the second case, one can speak about stochastic algorithms:
  - e.g., Monte Carlo EM (MCEM)
  - Stochastic EM (SEM)
  - Simulated Annealing EM (SAEM)

# Classification EM (CEM) algorithm

- we saw that EM computes the maximum likelihood (ML) estimate of a mixture model.
- The Classification EM (CEM) algorithm Celeux and Govaert (1992) estimates both the mixture model parameters and the classes' labels by maximizing the completed-data log-likelihood  $\mathcal{L}_c(\Psi; \mathbf{X}, \mathbf{z}) = \log p(\mathbf{X}, \mathbf{z}; \Psi)$
- start with an initial parameter  $\Psi^{(0)}$

- Step 1:** Compute the missing data  $\mathbf{z}^{(q+1)}$  given the observations and the current estimated model parameters  $\Psi^{(q)}$ :

$$\mathbf{z}^{(q+1)} = \arg \max_{\mathbf{z} \in \mathcal{Z}^n} \mathcal{L}_c(\Psi^{(q)}; \mathbf{X}, \mathbf{z})$$

- Step 2:** Compute the model parameters update  $\Psi^{(q+1)}$  by maximizing the complete-data log-likelihood given the current estimation of the missing data  $\mathbf{z}^{(q+1)}$ :

$$\Psi^{(q+1)} = \arg \max_{\Psi \in \Omega} \mathcal{L}_c(\Psi; \mathbf{X}, \mathbf{z}^{(q+1)}).$$



## Algorithm 2 Pseudo code of the CEM algorithm for GMMs.

**Inputs:** a data set  $\mathbf{X}$  and the number of clusters  $K$

fix a threshold  $\epsilon > 0$ ; set  $q \leftarrow 0$  (iteration)

**Initialize:**  $\Psi^{(0)} = (\pi_1^{(0)}, \dots, \pi_K^{(0)}, \Psi_1^{(0)}, \dots, \Psi_K^{(0)})$  with  $\Psi_k^{(0)} = (\mu_k^{(0)}, \Sigma_k^{(0)})$

**while** increment in the complete-data log-likelihood  $> \epsilon$  **do**

E-step:

**for**  $k = 1, \dots, K$  **do**

$$\text{Compute } \tau_{ik}^{(q)} = \frac{\pi_k \mathcal{N}(\mathbf{x}_i; \mu_k^{(q)}, \Sigma_k^{(q)})}{\sum_{\ell=1}^K \pi_\ell \mathcal{N}(\mathbf{x}_i; \mu_\ell^{(q)}, \Sigma_\ell^{(q)})}$$

**end for**

C-step:

**for**  $k = 1, \dots, K$  **do**

$$\text{Compute } z_i^{(q)} = \arg \max_{k \in \mathcal{Z}} \tau_{ik}^{(q)} \text{ for } i = 1, \dots, n$$

$$\text{Set } z_{ik}^{(q)} = 1 \text{ if } z_i^{(q)} = k \text{ and } z_{ik}^{(q)} = 0 \text{ otherwise, for } i = 1, \dots, n$$

**end for**

M-step:

**for**  $k = 1, \dots, K$  **do**

$$\text{Compute } \pi_k^{(q+1)} = \frac{\sum_{i=1}^n z_{ik}^{(q)}}{n}$$

$$\text{Compute } \mu_k^{(q+1)} = \frac{1}{n_k^{(q)}} \sum_{i=1}^n z_{ik}^{(q)} \mathbf{x}_i$$

$$\text{Compute } \Sigma_k^{(q+1)} = \frac{1}{n_k^{(q)}} \sum_{i=1}^n z_{ik}^{(q)} (\mathbf{x}_i - \mu_k^{(q+1)}) (\mathbf{x}_i - \mu_k^{(q+1)})^T$$

**end for**

$q \leftarrow q + 1$

**end while**

# CEM algorithm

- CEM is easy to implement, typically faster to converge than EM and monotonically improves the complete-data log-likelihood as the learning proceeds.
- converges toward a local maximum of the complete-data log-likelihood
- ! CEM provides biased estimates of the mixture model parameters. Indeed, CEM updates the model parameters from a truncated sample contrary to EM for which the model parameters are updated from the whole data through the fuzzy posterior probabilities and therefore the parameter estimations provided by EM are more accurate.
- **link with  $K$ -means:**
  - It can be shown that CEM which is formulated in a probabilistic framework, generalizes  $K$ -means
  - From a probabilistic point of view,  $K$ -means is equivalent to a particular case of the CEM algorithm for a mixture of  $K$  Gaussian densities with the same proportions  $\pi_k = \frac{1}{K} \forall k$  and identical isotropic covariance matrices  $\Sigma_k = \sigma^2 \mathbf{I} \forall k$ .

# Parsimonious Gaussian mixture models

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# Parsimonious Gaussian mixtures

- Parsimonious Gaussian mixture models<sup>1</sup> are statistical models that allow for capturing a specific cluster shapes (e.g., clusters having the same shape or different shapes, spherical or elliptical clusters, etc).
- Eigenvalue decomposition of the cluster covariance matrices:

$$\Sigma_k = \lambda_k \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k^T$$

where

- $\lambda_k$  represents the volume of the  $k$ th cluster (the amount of space of the cluster).
- $\mathbf{D}_k$  is a matrix with columns corresponding to the eigenvectors of  $\Sigma_k$  that determines the orientation of the cluster.
- $\mathbf{A}_k$  is a diagonal matrix, whose diagonal entries are the normalized eigenvalues of  $\Sigma_k$  arranged in a decreasing order and its determinant is 1. This matrix is associated with the shape of the cluster.

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<sup>1</sup>Banfield and Raftery (1993b); Celeux and Govaert (1995)

# Parsimonious Gaussian mixtures

- This eigenvalue decomposition provides three main families of models: the spherical family, the diagonal family, and the general family and produces 14 different models, according to the choice of the configuration for the parameters  $\lambda_k$ ,  $\mathbf{A}_k$ , and  $\mathbf{D}_k$

Decomposition	Model-Type	Prior	Applied to
$\lambda \mathbf{I}$	Spherical	$\mathcal{IG}$	$\lambda$
$\lambda_k \mathbf{I}$	Spherical	$\mathcal{IG}$	$\lambda_k$
$\lambda \mathbf{A}$	Diagonal	$\mathcal{IG}$	$\text{diag}(\lambda \mathbf{A})$
$\lambda_k \mathbf{A}$	Diagonal	$\mathcal{IG}$	$\text{diag}(\lambda_k \mathbf{A})$
$\lambda \mathbf{DAD}^T$	General	$\mathcal{IW}$	$\Sigma = \lambda \mathbf{DAD}^T$
$\lambda_k \mathbf{DAD}^T$	General	$\mathcal{IG}$ and $\mathcal{IW}$	$\lambda_k$ and $\Sigma = \mathbf{DAD}^T$
$\lambda_k \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k^T$	General	$\mathcal{IW}$	$\Sigma_k = \lambda_k \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k^T$

# Parsimonious GMMs

- In addition to providing flexible statistical models for the clusters, parsimonious Gaussian mixture can be viewed as techniques for reducing the number of parameters in the model.
- imposing constraints on the covariance matrices reduces the dimension of the optimization problem.
- The EM algorithms therefore provide more accurate estimations compared to the full mixture model.



# Model selection

- The complexity of a model  $\mathcal{M}$  is related to the number of its (free) parameters  $\nu$ , the penalty function then involves the number of model parameters.
- Let  $\mathcal{M}$  denote a model,  $\mathcal{L}(\boldsymbol{\theta})$  its log-likelihood and  $\nu$  the number of its free parameters. Consider that we fitted  $M$  different model structures  $(\mathcal{M}_1, \dots, \mathcal{M}_M)$ , from which we wish to choose the “best” one (ideally the one providing the best prediction on future data).
- Assume we have estimated the model parameters  $\hat{\boldsymbol{\theta}}_m$  for each model structure  $\mathcal{M}_m$  ( $m = 1, \dots, M$ ) from a sample of  $n$  observations  $\mathbf{X} = (\mathbf{x}_1, \dots, \mathbf{x}_n)$  and now we wish to choose among these fitted models.









# Examples

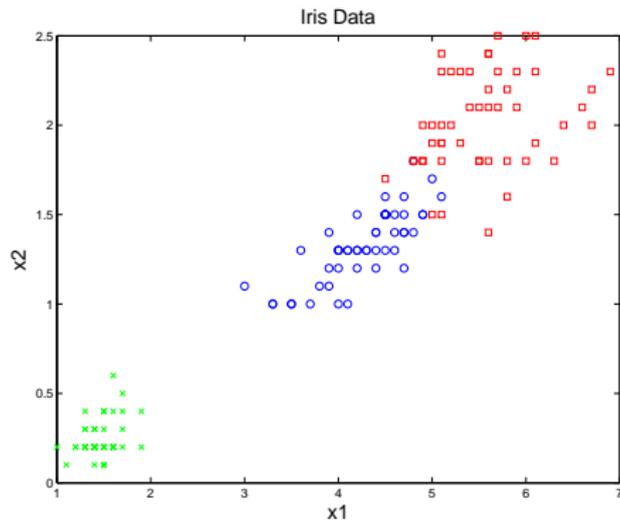


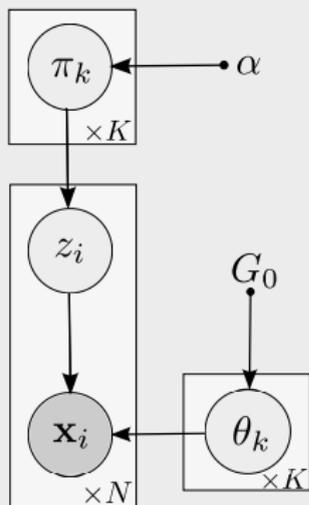
Figure : Iris data of Fisher: The data are colored according to the true partition.

# Bayesian regularization of mixtures and Model-Based Clustering

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# The Bayesian finite mixture model

Graphical model representation of the Bayesian finite mixture model

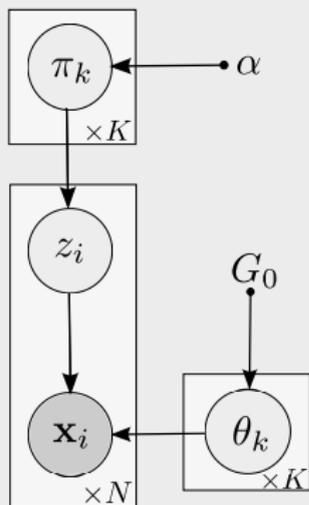


Generative model

$$\pi_1, \dots, \pi_K | \alpha \sim \text{Dir}(\alpha_1, \dots, \alpha_K)$$

# The Bayesian finite mixture model

Graphical model representation of the Bayesian finite mixture model



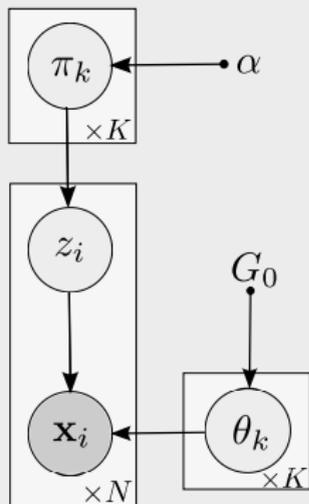
Generative model

$$\pi_1, \dots, \pi_K | \boldsymbol{\alpha} \sim \text{Dir}(\alpha_1, \dots, \alpha_K)$$

$$z_i | \boldsymbol{\pi} \sim \text{Mult}(\cdot | \boldsymbol{\pi})$$

# The Bayesian finite mixture model

Graphical model representation of the Bayesian finite mixture model



Generative model

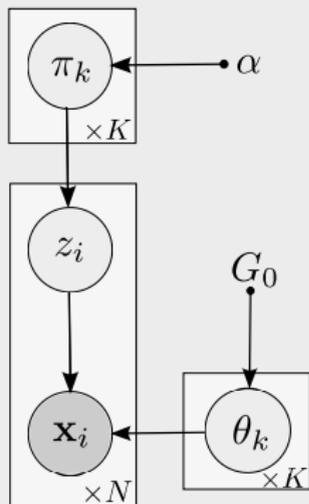
$$\pi_1, \dots, \pi_K | \boldsymbol{\alpha} \sim \text{Dir} | (\alpha_1, \dots, \alpha_K)$$

$$z_i | \boldsymbol{\pi} \sim \text{Mult}(\cdot | \boldsymbol{\pi})$$

$$\boldsymbol{\theta}_{z_i} | G_0 \sim G(\cdot | G_0)$$

# The Bayesian finite mixture model

Graphical model representation of the Bayesian finite mixture model



## Generative model

$$\pi_1, \dots, \pi_K | \boldsymbol{\alpha} \sim \text{Dir} | (\alpha_1, \dots, \alpha_K)$$

$$z_i | \boldsymbol{\pi} \sim \text{Mult}(\cdot | \boldsymbol{\pi})$$

$$\boldsymbol{\theta}_{z_i} | G_0 \sim G(\cdot | G_0)$$

$$\mathbf{x}_i | z_i, \boldsymbol{\theta}_{z_i} \sim f(\cdot | \boldsymbol{\theta}_{z_i})$$

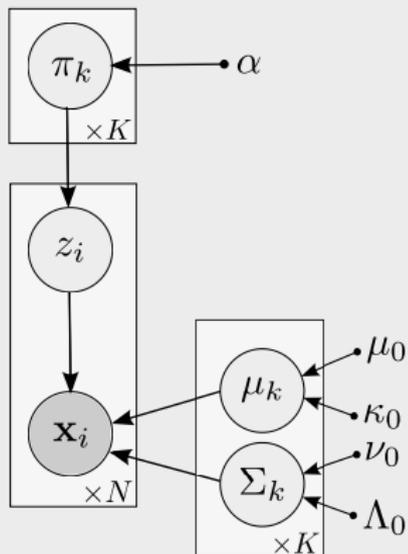
$$\boldsymbol{\theta}_{z_i} = (\boldsymbol{\mu}_{z_i}, \boldsymbol{\Sigma}_{z_i})$$

G : prior distribution

$G_0$  : hyperparameters

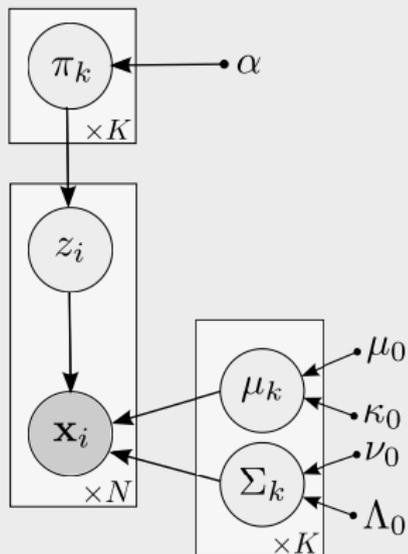
# The Bayesian finite GMM

Graphical model representation of the Bayesian finite GMM



# The Bayesian finite GMM

Graphical model representation of the Bayesian finite GMM

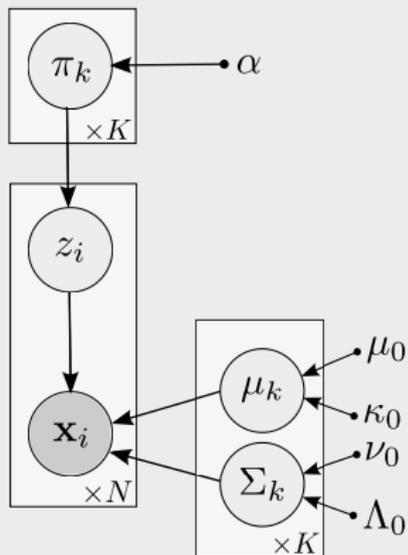


Generative model

$$\pi_1, \dots, \pi_K | \alpha \sim \text{Dir} | (\alpha_1, \dots, \alpha_K)$$

# The Bayesian finite GMM

Graphical model representation of the Bayesian finite GMM



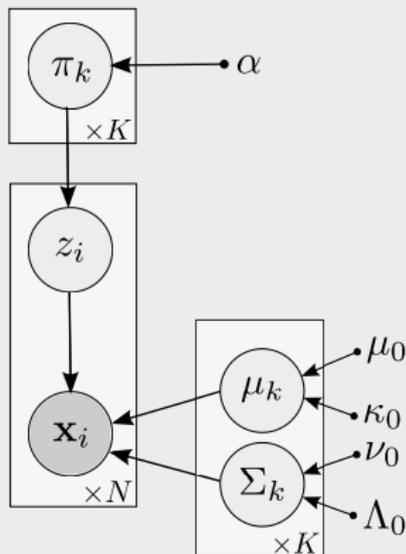
Generative model

$$\pi_1, \dots, \pi_K | \boldsymbol{\alpha} \sim \text{Dir} | (\alpha_1, \dots, \alpha_K)$$

$$z_i | \boldsymbol{\pi} \sim \text{Mult}(\cdot | \boldsymbol{\pi})$$

# The Bayesian finite GMM

Graphical model representation of the Bayesian finite GMM



## Generative model

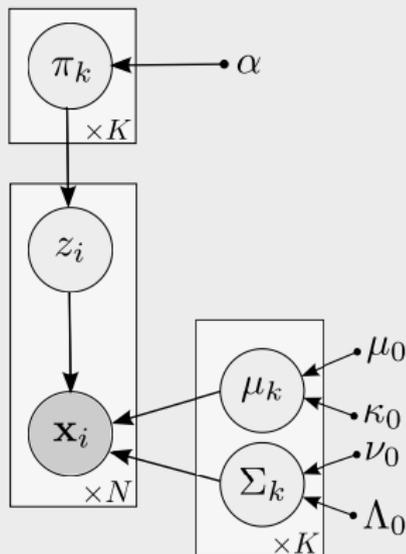
$$\pi_1, \dots, \pi_K | \alpha \sim \text{Dir} | (\alpha_1, \dots, \alpha_K)$$

$$z_i | \pi \sim \text{Mult}(. | \pi)$$

$$\theta_{z_i} | G_0 \sim G(. | G_0)$$

# The Bayesian finite GMM

Graphical model representation of the Bayesian finite GMM



## Generative model

$$\pi_1, \dots, \pi_K | \alpha \sim \text{Dir} | (\alpha_1, \dots, \alpha_K)$$

$$z_i | \boldsymbol{\pi} \sim \text{Mult}(\cdot | \boldsymbol{\pi})$$

$$\boldsymbol{\theta}_{z_i} | G_0 \sim G(\cdot | G_0)$$

$$\mathbf{x}_i | z_i, \boldsymbol{\theta}_{z_i} \sim f(\cdot | \boldsymbol{\theta}_{z_i})$$

$$\boldsymbol{\theta}_k = (\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)$$

$$\boldsymbol{\mu}_k \sim \mathcal{N}(\boldsymbol{\mu}_0, \mathbf{V}_0)$$

$$\boldsymbol{\Sigma}_k \sim \mathcal{IW}(\mathbf{S}_0, \nu_0)$$

# The Gibbs sampler for the Bayesian finite GMM

## Bayesian sampling

$$\begin{aligned}
 z_i | \mathbf{X}, \boldsymbol{\mu}, \boldsymbol{\theta} &\sim \text{Mult}(\cdot | \tau_{i1}, \dots, \tau_{iK}) \\
 \tau_{ik} &= p(z_i = k | \mathbf{x}_i, \boldsymbol{\pi}, \boldsymbol{\theta}) = \frac{\pi_k \mathcal{N}(\mathbf{x}_i | \boldsymbol{\theta}_k)}{\sum_{l=1}^K \pi_l \mathcal{N}(\mathbf{x}_i | \boldsymbol{\theta}_l)}
 \end{aligned} \tag{1}$$

# The Gibbs sampler for the Bayesian finite GMM

## Bayesian sampling

$$z_i | \mathbf{X}, \boldsymbol{\mu}, \boldsymbol{\theta} \sim \text{Mult}(\cdot | \tau_{i1}, \dots, \tau_{iK}) \quad (1)$$

$$\tau_{ik} = p(z_i = k | \mathbf{x}_i, \boldsymbol{\pi}, \boldsymbol{\theta}) = \frac{\pi_k \mathcal{N}(\mathbf{x}_i | \boldsymbol{\theta}_k)}{\sum_{l=1}^K \pi_l \mathcal{N}(\mathbf{x}_i | \boldsymbol{\theta}_l)}$$

$$\pi_1, \dots, \pi_K | \mathbf{z} \sim \text{Dir}(\cdot | \alpha_1 + n_1, \dots, \alpha_K + n_K) \quad (2)$$

$$n_k = \sum_{i=1}^n z_{ik}$$

# The Gibbs sampler for the Bayesian finite GMM

## Bayesian sampling

$$z_i | \mathbf{X}, \boldsymbol{\mu}, \boldsymbol{\theta} \sim \text{Mult}(\cdot | \tau_{i1}, \dots, \tau_{iK}) \quad (1)$$

$$\tau_{ik} = p(z_i = k | \mathbf{x}_i, \boldsymbol{\pi}, \boldsymbol{\theta}) = \frac{\pi_k \mathcal{N}(\mathbf{x}_i | \boldsymbol{\theta}_k)}{\sum_{l=1}^K \pi_l \mathcal{N}(\mathbf{x}_i | \boldsymbol{\theta}_l)}$$

$$\pi_1, \dots, \pi_K | \mathbf{z} \sim \text{Dir}(\cdot | \alpha_1 + n_1, \dots, \alpha_K + n_K) \quad (2)$$

$$n_k = \sum_{i=1}^n z_{ik}$$

$$\boldsymbol{\mu}_k | \boldsymbol{\Sigma}_k, \mathbf{z}, \mathbf{X} \sim \mathcal{N}(\cdot | \mathbf{m}_k, \mathbf{V}_k) \quad (3)$$

$$\mathbf{V}_k^{-1} = \mathbf{V}_0^{-1} + n_k \boldsymbol{\Sigma}_k^{-1}$$

$$\mathbf{m}_k = \mathbf{V}_k (\boldsymbol{\Sigma}_k^{-1} \sum_{i=1}^n z_{ik} \mathbf{x}_i + \mathbf{V}_0^{-1} \mathbf{m}_0)$$

# The Gibbs sampler for the Bayesian finite GMM

## Bayesian sampling

$$z_i | \mathbf{X}, \boldsymbol{\mu}, \boldsymbol{\theta} \sim \text{Mult}(\cdot | \tau_{i1}, \dots, \tau_{iK}) \quad (1)$$

$$\tau_{ik} = p(z_i = k | \mathbf{x}_i, \boldsymbol{\pi}, \boldsymbol{\theta}) = \frac{\pi_k \mathcal{N}(\mathbf{x}_i | \boldsymbol{\theta}_k)}{\sum_{l=1}^K \pi_l \mathcal{N}(\mathbf{x}_i | \boldsymbol{\theta}_l)}$$

$$\pi_1, \dots, \pi_K | \mathbf{z} \sim \text{Dir}(\cdot | \alpha_1 + n_1, \dots, \alpha_K + n_K) \quad (2)$$

$$n_k = \sum_{i=1}^n z_{ik}$$

$$\boldsymbol{\mu}_k | \boldsymbol{\Sigma}_k, \mathbf{z}, \mathbf{X} \sim \mathcal{N}(\cdot | \mathbf{m}_k, \mathbf{V}_k) \quad (3)$$

$$\mathbf{V}_k^{-1} = \mathbf{V}_0^{-1} + n_k \boldsymbol{\Sigma}_k^{-1}$$

$$\mathbf{m}_k = \mathbf{V}_k (\boldsymbol{\Sigma}_k^{-1} \sum_{i=1}^n z_{ik} \mathbf{x}_i + \mathbf{V}_0^{-1} \mathbf{m}_0)$$

$$\boldsymbol{\Sigma}_k | \boldsymbol{\mu}_k, \mathbf{z}, \mathbf{X} \sim \mathcal{IW}(\mathbf{S}_0 + \sum_{i=1}^n z_{ik} (\mathbf{x}_i - \boldsymbol{\mu}_k)(\mathbf{x}_i - \boldsymbol{\mu}_k)^T, \nu_0 + n_k) \quad (4)$$

# Infinite Gaussian Mixture Model and Dirichlet Process Mixtures

- Infinite GMM:  $p(\mathbf{x}_i|\boldsymbol{\theta}) = \sum_{k=1}^{\infty} \pi_k \mathcal{N}_k(\mathbf{x}_i|\boldsymbol{\theta}_k)$  Rasmussen (2000)
- Parameters:  $\boldsymbol{\theta} = \{\pi_k, \boldsymbol{\theta}_k = (\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)\}_{k=1}^{\infty}$
- Prior: add a distribution over the parameters distribution: a Dirichlet Process Antoniak (1974)
- Generative model:

$$G|\alpha, G_0 \sim \text{DP}(\alpha, G_0)$$

$$\boldsymbol{\theta}_i|G \sim G$$

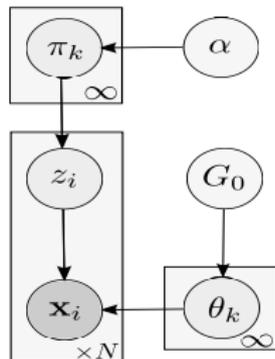
$$\mathbf{x}_i|\boldsymbol{\theta}_i \sim p(\cdot|\boldsymbol{\theta}_i)$$

equivalent to

$$z_i|\alpha \sim \text{CRP}(z_i; \alpha)$$

$$\boldsymbol{\theta}_{z_i}|G_0 \sim G_0$$

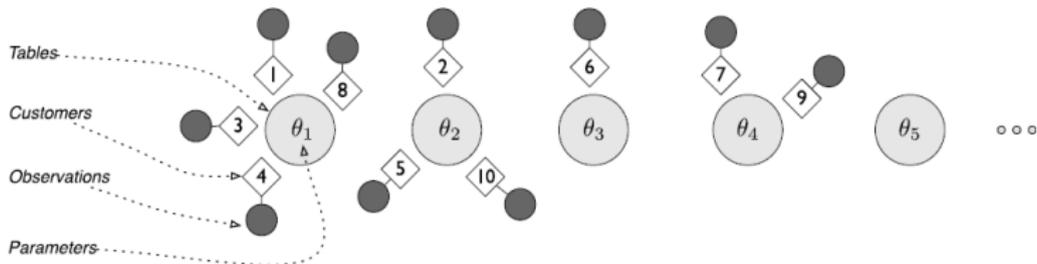
$$\mathbf{x}_i|\boldsymbol{\theta}_{z_i} \sim p(\cdot|\boldsymbol{\theta}_{z_i})$$



# Chinese Restaurant Process (CRP)

Imagine a Restaurant with an infinite number of tables and in which customers are entering and sitting at these tables.

- ① The first customer sits at table 1
- ② The second customer may sit at table with probability  $\frac{1}{1+\alpha}$  or chose another table with probability  $\frac{\alpha}{1+\alpha}$
- ③ ...
- ④  $i$ th customer sits at table  $k$  with probability proportional to the number of already seated customers  $n_k$  and may choose a new table with a probability proportional to a small positive real number  $\alpha$



# Chinese Restaurant Process (CRP)

Chinese Restaurant Process (CRP) Wood and Black (2008); Samuel and Blei (2012):

- The CRP provides a distribution on the infinite partitions of the data:  
 $p(\mathbf{z}) = p(z_1)p(z_2|z_1)\dots p(z_n|z_{n-1})$ :

$$\begin{aligned}
 p(z_i = k | z_1, \dots, z_{i-1}) &= \text{CRP}(z_1, \dots, z_{i-1}; \alpha) \\
 &= \begin{cases} \frac{n_k}{i-1+\alpha} & \text{if } k \leq K_+ \\ \frac{\alpha}{i-1+\alpha} & \text{if } k > K_+ \end{cases}
 \end{aligned}$$

- $\alpha$  represents the CRP concentration parameter
- $K_+$  : nbr. of tables for which the nbr. of customers sitting in is  $n_k > 0$  (active clusters)
- $k \leq K_+$  means that  $k$  is a previously occupied table and  $k > K_+$  means  $k$  is a new table to be occupied.

# Gibbs sampler for the Infinite Parsimonious GMM

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## Algorithm 3 Gibbs sampler for the Infinite Parsimonious GMM

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**Entrees:** Data  $\mathbf{x}_i$ , nbr of Gibbs samples  $n_S$ .

Initialization:  $q \leftarrow 0$ ; hyper-parameters  $G_0^{(q)}$ ,  $\alpha$ ; nbr of clusters  $K_+ = 1$ .

**for**  $i = 1, \dots, n$  **do**

    sample the cluster labels  $z_i^{(t)} \sim p(\mathbf{x}_i | z_i, \theta_k)$  CRP( $\{z_1, \dots, z_n\} \setminus z_i; \alpha^{(t)}$ )

    if  $z_i^{(t)} = K_+ + 1$  create a new cluster  $K_+ = K_+ + 1$ , sample  $\theta_{z_i}^{(t)}$  from the prior

**end for**

**for**  $k = 1, \dots, K_+$  **do**

    sample  $\theta_k^{(t)}$  from the posterior

**end for**

sample  $\alpha^{(t)}$

$\mathbf{z}^{(t+1)} \leftarrow \mathbf{z}^{(t)}$

$\alpha^{(t+1)} \leftarrow \alpha^{(t)}$

$t \leftarrow t + 1$

**Outputs:**  $\{\hat{\theta}, \hat{z}, \hat{K} = K_+\}$

Seven included models in the non-parametric approach:

Decomposition	Model-Type	Prior	Applied to
$\lambda \mathbf{I}$	Spherical	$\mathcal{I}\mathcal{G}$	$\lambda$
$\lambda_k \mathbf{I}$	Spherical	$\mathcal{I}\mathcal{G}$	$\lambda_k$
$\lambda \mathbf{A}$	Diagonal	$\mathcal{I}\mathcal{G}$	$\text{diag}(\lambda \mathbf{A})$
$\lambda_k \mathbf{A}$	Diagonal	$\mathcal{I}\mathcal{G}$	$\text{diag}(\lambda_k \mathbf{A})$
$\lambda \mathbf{DAD}^T$	General	$\mathcal{I}\mathcal{W}$	$\Sigma = \lambda \mathbf{DAD}^T$
$\lambda_k \mathbf{DAD}^T$	General	$\mathcal{I}\mathcal{G}$ and $\mathcal{I}\mathcal{W}$	$\lambda_k$ and $\Sigma = \mathbf{DAD}^T$
$\lambda_k \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k^T$	General	$\mathcal{I}\mathcal{W}$	$\Sigma_k = \lambda_k \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k^T$

## Bayesian learning

Gibbs sampler: e.g for  $\lambda_k \mathbf{DAD}^T$ : Normal Inverse-Wishart (prior/posterior)

$$\mu_k | \cdot \sim \mathcal{N}(\mu_0, \lambda_k \Sigma_0 / \kappa_0) \dots \Sigma_0 | \cdot \sim \mathcal{I}\mathcal{W}(v_0, \Lambda_0) \dots \lambda_k | \cdot \sim \mathcal{I}\mathcal{G}(r_0/2, s_0/2)$$

$$\mu_k | \mathbf{X}_{\cdot,} \sim \mathcal{N}\left(\frac{n_k \bar{\mathbf{x}}_k + \kappa_0 \mu_0}{n_k + \kappa_0}, \frac{\lambda_k \Sigma_0}{n_k + \kappa_0}\right)$$

$$\Sigma_0 | \mathbf{X}_{\cdot,} \sim \mathcal{I}\mathcal{W}\left(v_0 + n, \Lambda_0 + \sum_{k=1}^K \left\{ \frac{W_k}{\lambda_k} + \frac{n_k \kappa_0}{\lambda_k (n_k + \kappa_0)} (\bar{\mathbf{x}}_k - \mu_0)(\bar{\mathbf{x}}_k - \mu_0)^T \right\}\right)$$

$$\lambda_k | \mathbf{X}_{\cdot,} \sim \mathcal{I}\mathcal{G}\left(\frac{r_0 + n_k d}{2}, \frac{1}{2} \left\{ s_0 + \text{tr}(W_k \Sigma_0^{-1}) + \frac{n_k \kappa_0}{n_k + \kappa_0} (\bar{\mathbf{x}}_k - \mu_0)^T \Sigma_0^{-1} (\bar{\mathbf{x}}_k - \mu_0) \right\}\right)$$

# Markovian Extension

The infinite GMMs has been extended to the infinite HMM for sequential data modeling: This is the Hierarchical Dirichlet Process for Hidden Markov Model (HDP-HMM)

Two main inference approaches approaches: the Gibbs sampling <sup>2 3</sup> and the Beam sampling <sup>4</sup>.

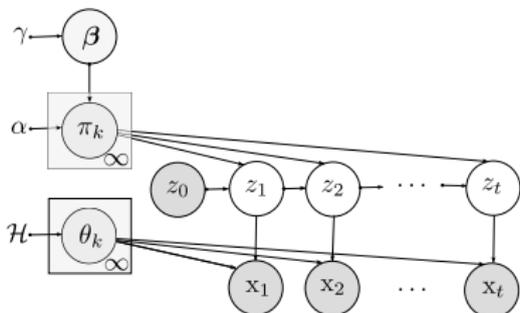


Figure : Infinite Hidden Markov Model (IHMM) graphical representation

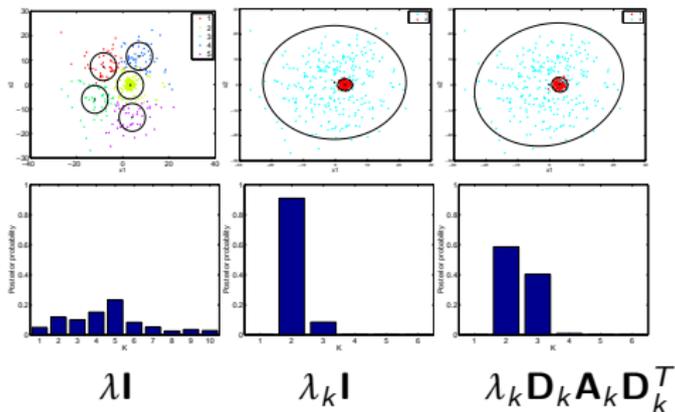
<sup>2</sup>Fox et al. (2008)

<sup>3</sup>Teh et al. (2006)

<sup>4</sup>Van Gael et al. (2008)

# Experiment on simulated data

- 1 A two-clusters data set
- 2  $n = 500$  observation in  $\mathbb{R}^2$
- 3  $\boldsymbol{\pi} = [.5 \ .5]$ ;  $\boldsymbol{\mu}_1 = [0 \ 0]^T$ ;  $\boldsymbol{\mu}_2 = [3 \ 0]^T$ ;  $\boldsymbol{\Sigma}_1 = 100 * \mathbf{I}_2$ ;  $\boldsymbol{\Sigma}_2 = \mathbf{I}_2$



# Experiments on benchmarks

Dataset	$n$	$d$	True $K$
Iris	150	4	3
Old Faithful Geyser	272	2	?
Trees	31	3	?
Wine	178	13	3
Diabetes	145	3	3

Model	Iris	Geyser	Trees	Wine	Diabetes
$\lambda \mathbf{I}$	4	2	1	1	3
$\lambda_k \mathbf{I}$	3	2	1	2	5
$\lambda \mathbf{A}$	3	3	2	3	3
$\lambda_k \mathbf{A}$	3	2	2	1	5
$\lambda \mathbf{DAD}^T$	4	2	2	3	5
$\lambda_k \mathbf{DAD}^T$	2	2	2	3	3
$\lambda_k \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k^T$	2	2	2	3	3

Table : Estimated  $K$  by the infinite parsimonious GMM

# Experiments on Benchmarks

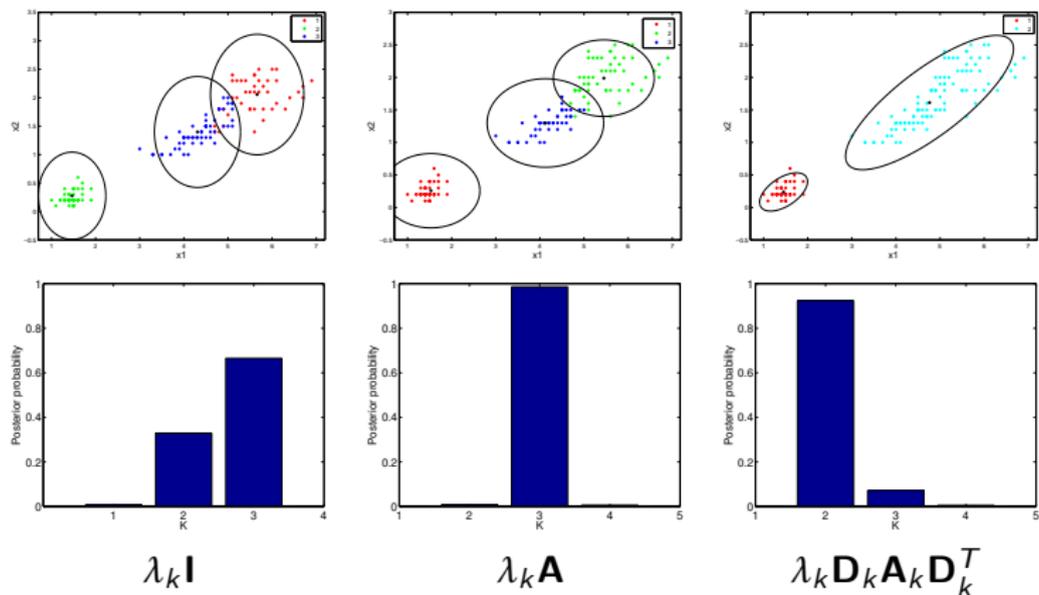


Figure : Graphical results for Iris data

# Experiments on Benchmarks

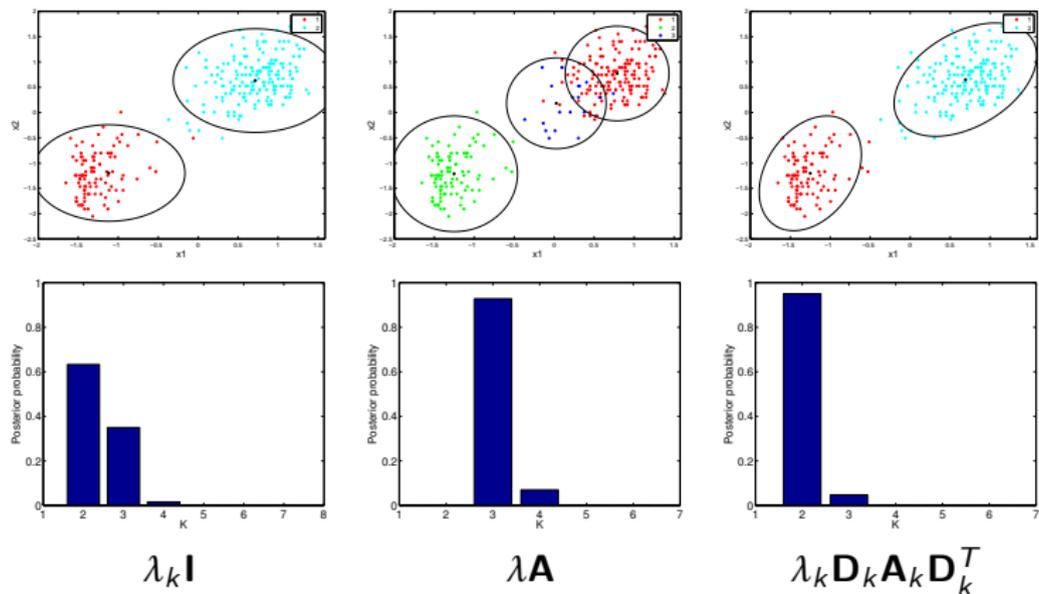


Figure : Graphical results for Old Faithful Geyser data

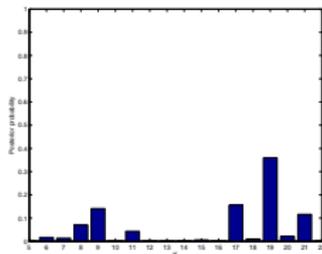
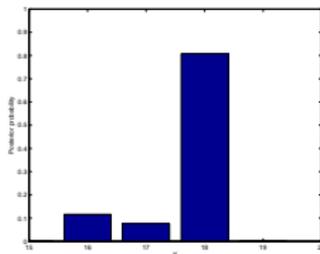
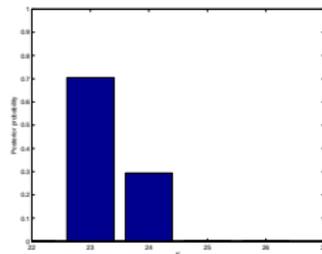
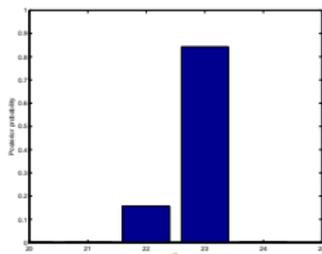
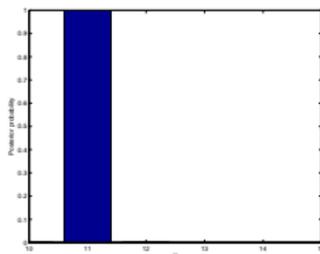
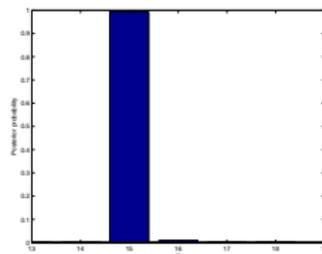
# Whale song decomposition

- In this experiment, we apply the proposed approach to a challenging problem of humpback whale song decomposition.
- The analysis is unsupervised and aims at discovering the call units (which can be considered as a kind of whale vocabulary),
- this can be seen as a problem of unsupervised call units classification as in<sup>5</sup>.
- We therefore reformulate the problem of whale song decomposition as a clustering problem.
- We apply our proposed IPGMM to find a partition of the whale song into clusters, and automatically infer the number of clusters from the data.
- We then extend the Infinite GMM to the Markovian case as in
- The data consist of MFCC parameters of 8.6 minutes of a Humpback whale song recordings

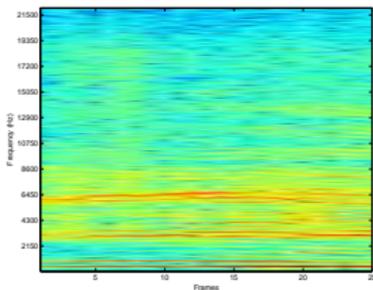
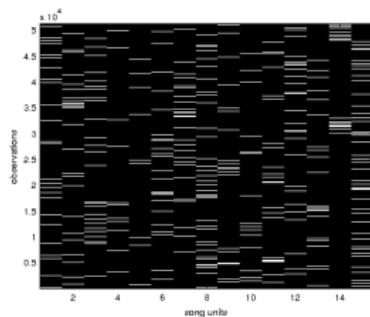
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<sup>5</sup>Pace et al. (2010)

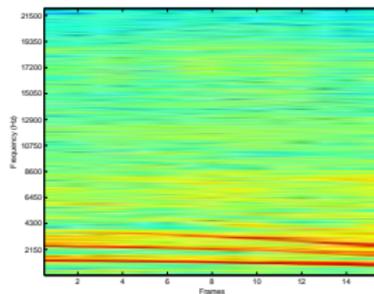
# whale song decomposition problem

 $\lambda_I$  $\lambda_A$  $\lambda_k \mathbf{D} \mathbf{A} \mathbf{D}^T$  $\lambda_k \mathbf{I}$  $\lambda_k \mathbf{A}$  $\lambda_k \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k^T$

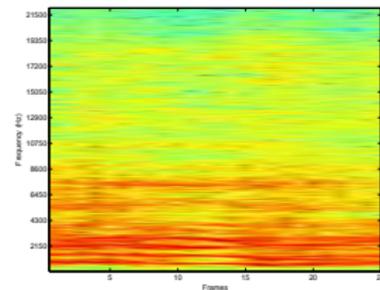
# whale song decomposition problem



song unit 1



song unit 9



song unit 14

State sequence obtained by using a HDP-HMM <sup>6</sup>.

"sparse" approach: the number of estimated states equals  $K = 6$

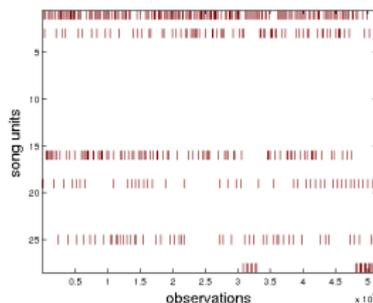
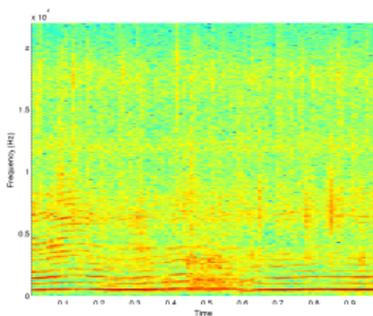
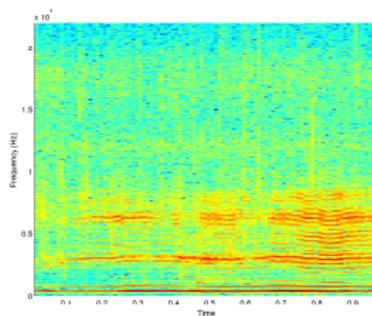


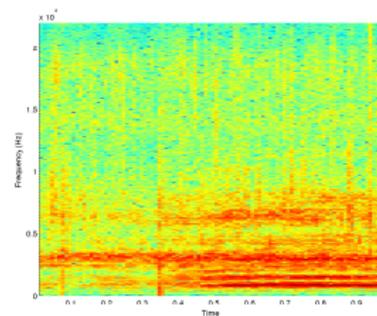
Figure : State sequence obtained by an Infinite HMM.



song unit 16



song unit 19



song unit 30

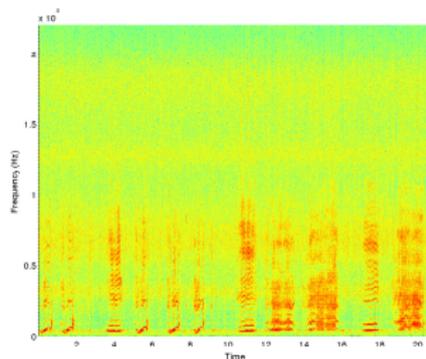


Figure : The state sequences (left) and the spectrogram of the ninth unit (right) obtained by the HDP-HMM (Beam sampling)

# Conclusion and Perspectives

- Mixtures are very flexible for cluster analysis namely via Parsimonious mixture modeling
- The Dirichlet Process mixture approach is a Bayesian non-parametric alternative
- It avoids the problem of model selection encountered in maximum likelihood and Bayesian learning of parametric GMMs
- 
- The parsimonious version allows to have several flexible models adapted for several clusters configurations
- Perspectives :
  - More comparisons between the different modes (e.g. using Bayes Factors)
  - More experiments for the Markovian extension for sequential data
  - Infinite block mixture
  - Other MCMC techniques

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Thank you!