

Data Science

Static data analysis

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Master en Sciences Informatiques - Semestre d'Automne

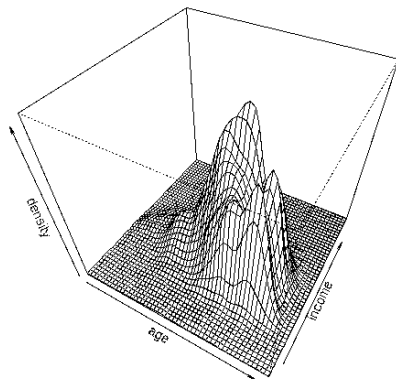
Data modelling

- ★ Up until now, we have studied data with an implicit model underlying the technique
 - Component models → Variance as a criterion on centered data (Normal distribution)
 - Discriminant models → Variance of projected data in within- and between-class models
- ⇒ The distribution is fixed (essentially normal) and we look for its parameters (μ , σ)
- ★ Alternatively we can search a model for data density
- ★ Let $f(\mathbf{x}) : \mathcal{F} \rightarrow \mathbb{R}$ be the data density

Density estimation

- ★ Nearest neighbor methods (knn)
- ★ Parzen windows, RBF networks
- ★ Histograms
- ★ Mixture models

Density estimation: perspective plot



Mixture models

Definition

- ★ The density $f(x)$ is generated by c “basis” functions (components)

$$f(x) = \sum_{j=1}^c \pi_j \phi(x, \theta_j)$$

- ★ π_j are the mixture parameters
- ★ $\phi(x, \theta_j)$ are functions controlled by parameters θ_j

Hypotheses

1. The number of components (c) is known
2. The family of functions ϕ is known
3. Labels (classes) are unknown

Probabilist reading

- ★ Density $f(x)$ represents a random process where x is drawn from a set of states ω_j with prior probability $P(\omega_j)$

$$f(x) = \sum_{j=1}^c \pi_j \phi(x, \theta_j)$$

$$f(x) = p(x|\theta) = \sum_{j=1}^c P(\omega_j) P(x|\omega_j, \theta_j)$$

- ★ We get $\pi_j = P(\omega_j)$ and $\sum_j \pi_j = 1$
- ★ $\phi(x, \theta_j)$ is the conditional probability that x is generated by ω_j

Gaussian mixture

ϕ is a probability density function. Choosing the (agnostic) normal law as basis $\mathcal{N}(\mu, \Sigma)$ seems reasonable:

$$f(x) = \sum_{j=1}^c \pi_j \mathcal{N}(\mu_j, \Sigma_j)$$

- ★ Can approximate any density
- ★ Enables a linear system of its parameters by maximising the log-likelihood

Maximum log-likelihood (ML)

- ★ Given $\Omega = \{x_1, \dots, x_N\}$ unlabeled samples generated by the mixture $f(x) = p(x|\theta)$.
- ★ $\theta = \{\pi_j, \mu_j, \Sigma_j\}$ are the parameters to infer.
- ★ Likelihood :

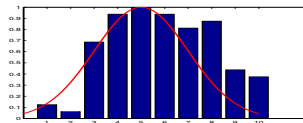
$$p(\Omega|\theta) = \prod_i^N p(x_i|\theta)$$

- ★ Estimation : $\hat{\theta} = \arg \max_{\theta} p(\Omega|\theta)$
- ★ or maximum **log-likelihood**

$$l(\theta, \Omega) = \sum_{i=1}^N \log p(x_i|\theta) = \sum_{i=1}^N \log \left[\sum_{j=1}^c \pi_j \phi(x_i, \theta_j) \right]$$

Basic case : 1 component, $c = 1$

$$\star \theta = \{\mu, \Sigma\}$$



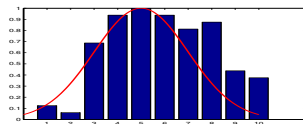
$$\max_{\theta} \sum_i \log e^{-(x_i - \mu)^T \Sigma^{-1} (x_i - \mu)}$$



$$\min_{\theta} \sum_i (x_i - \mu)^T \Sigma^{-1} (x_i - \mu)$$

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$$\max_{\theta} \sum_i \log e^{-(x_i - \mu)^T \Sigma^{-1} (x_i - \mu)}$$



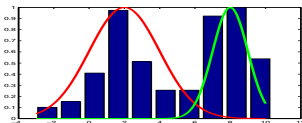
$$\min_{\theta} \sum_i (x_i - \mu)^T \Sigma^{-1} (x_i - \mu)$$

$$\star \hat{\mu} = \frac{1}{N} \sum_i x_i$$

$$\star \hat{\Sigma} = \frac{1}{N} \sum_i (x_i - \mu)(x_i - \mu)^T$$

A bit more complex: 2 components

$$\begin{aligned}\theta &= \{\pi, \theta_1, \theta_2\} \\ &= \{\pi_1, \pi_2, \mu_1, \Sigma_1, \mu_2, \Sigma_2\}\end{aligned}$$



$$l(\theta, \Omega) = \sum_i^N \log [(1 - \pi)\phi(x_i, \theta_1) + \pi\phi(x_i, \theta_2)]$$

- ⇒ difficult to maximise because of the sum inside the **log**!
- ⇒ Solution : iteratif 2 steps (E-M) algorithm to maximise /
- **Expectation-Maximisation** (EM) algorithm

EM algorithm (2 components)

- ★ The unknown here is the assignement x_i to one of the 2 components ϕ_i
- ⇒ If we knew it, we would treat the problem as twice 1 component
- ★ The EM algorithm introduces **unknown variables** : the assignement $\Delta_i \in \{0, 1\}$ of every x_i to one component :

$$x_i \sim \phi_1 \text{ if } \Delta_i = 0, x_i \sim \phi_2 \text{ if } \Delta_i = 1$$

Expectation (E-step)

- ★ Assume we know an initial value for θ^0
- ★ We can infer the contribution of every data x_i to every density (parameterized by θ_i^0):

$$\begin{aligned}\gamma_i(\theta^0) &= E[\Delta_i | \theta^0, \Omega] \\ &= \frac{\pi \phi(x_i, \theta_2^0)}{(1 - \pi) \phi(x_i, \theta_1^0) + \pi \phi(x_i, \theta_2^0)}\end{aligned}$$

- ★ γ_i is the **responsability**.
- ★ It is the **expectation** of Δ_i over all components

Responsability and soft-assignment

γ_i allows to determine $\Delta_i \Rightarrow x_i$ can be assigned to either ϕ_1 or ϕ_2
 \Rightarrow K-means-type **hard-assignment**. Each data is assigned to one and only one cluster

EM is “softer”. A data may contribute (via γ_i) to several density modes (clusters). EM computes a **soft-assignment** (with $\sum_i \gamma_i = 1$)

Maximisation (M-step)

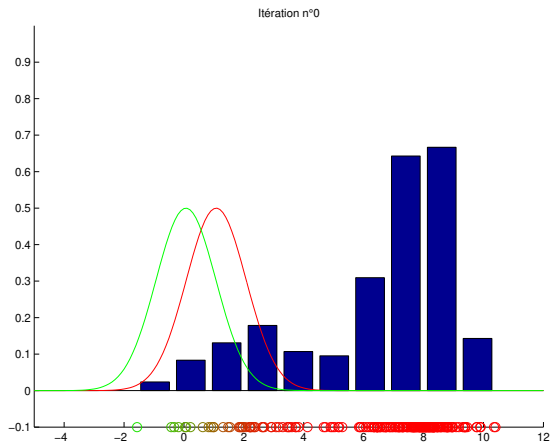
- ★ Given every data responsibility (γ_i), we can estimate the parameters by (weighted) maximum likelihood:

$$\begin{aligned}\hat{\mu}_1 &= \frac{\sum_{i=1}^N (1 - \gamma_i) x_i}{\sum_{i=1}^N 1 - \gamma_i} & \hat{\Sigma}_1 &= \frac{\sum_{i=1}^N (1 - \gamma_i) (x_i - \mu_1)(x_i - \mu_1)^T}{\sum_{i=1}^N 1 - \gamma_i} \\ \hat{\mu}_2 &= \frac{\sum_{i=1}^N \gamma_i x_i}{\sum_{i=1}^N \gamma_i} & \hat{\Sigma}_2 &= \frac{\sum_{i=1}^N \gamma_i (x_i - \mu_1)(x_i - \mu_1)^T}{\sum_{i=1}^N \gamma_i}\end{aligned}$$

- ★ Proportion for mixture 1: $\pi = \sum_{i=1}^N \gamma_i / N$

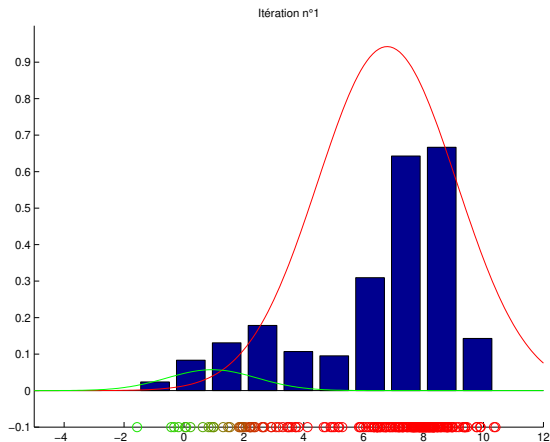
Illustration

Successive iterations of E- and M-steps



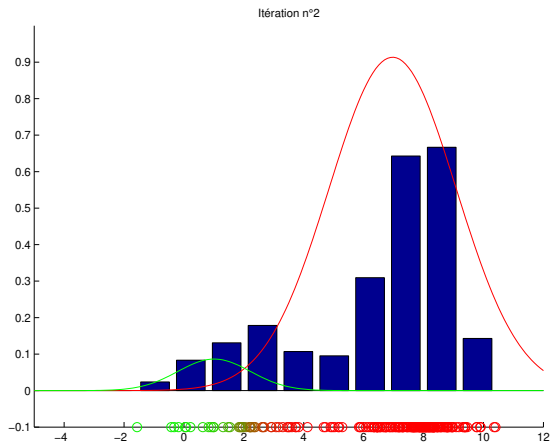
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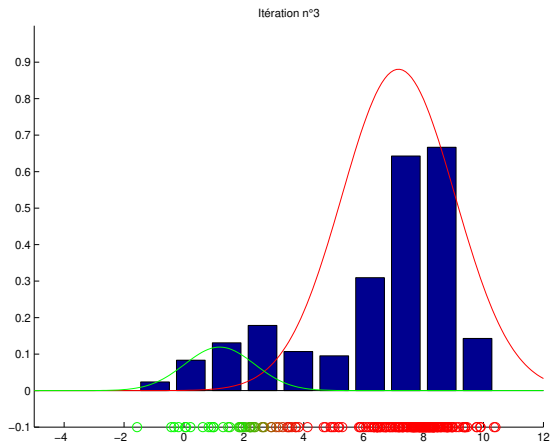
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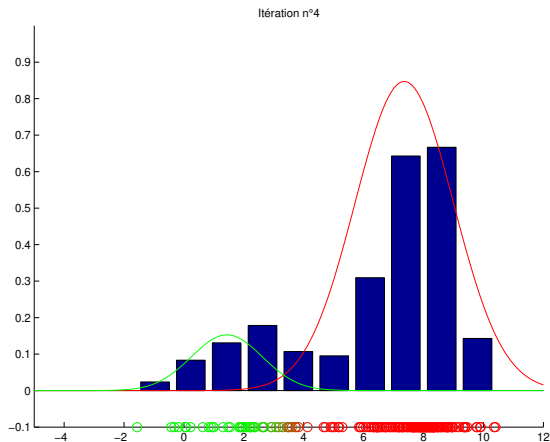
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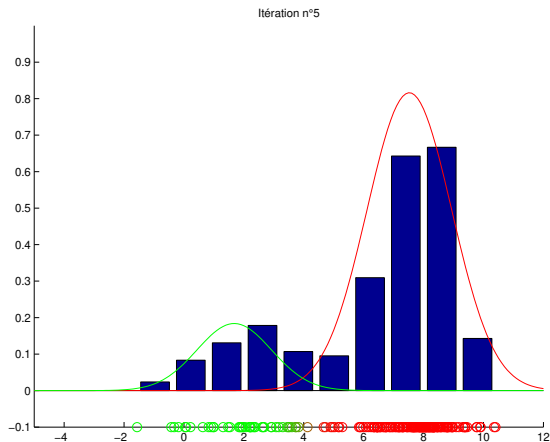
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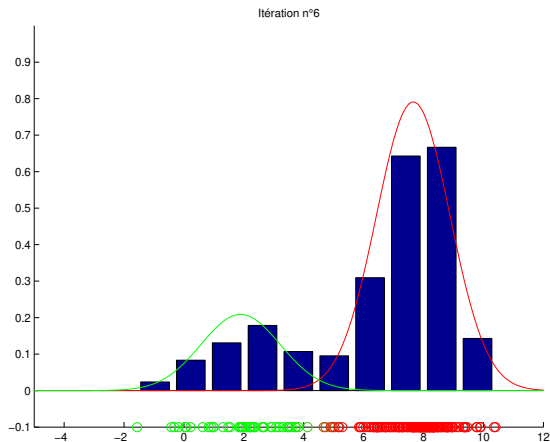
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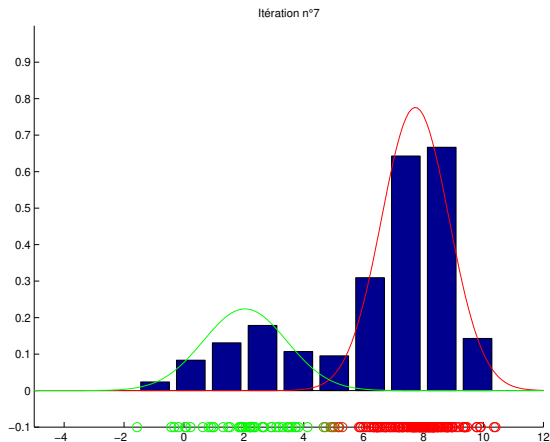
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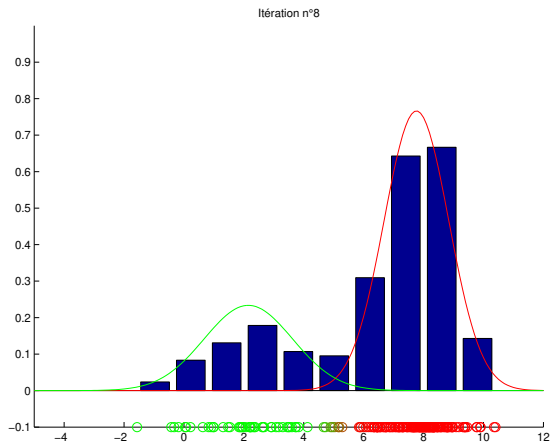
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Successive iterations of E- and M-steps



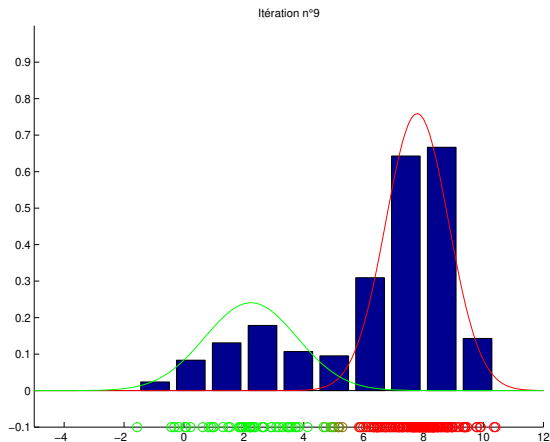
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Successive iterations of E- and M-steps



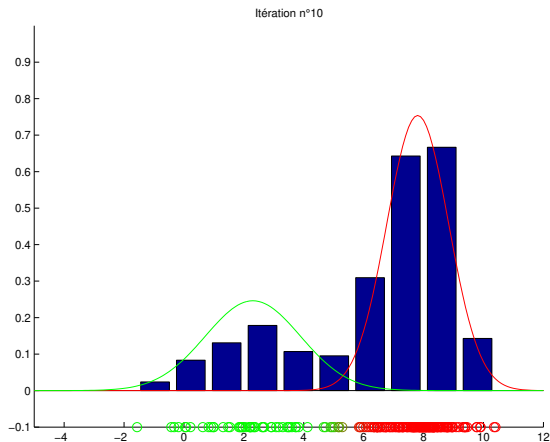
Illustration

Successive iterations of E- and M-steps



Illustration

Successive iterations of E- and M-steps



Results

★ True parameters

π	μ_1	σ_1	μ_2	σ_2
0.75	2	2	8	1

★ Estimated parameters

$\hat{\pi}$	$\hat{\mu}_1$	$\hat{\sigma}_1$	$\hat{\mu}_2$	$\hat{\sigma}_2$
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10 iterations

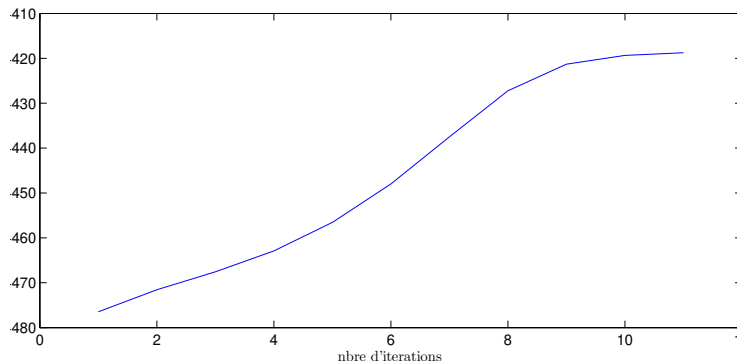
0.76	2.17	1.56	7.91	1.06
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20 iterations

0.76	2.15	1.98	8.01	0.98
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Iterations

Alternate cycle Expectation-Maximisation → increases the likelihood of the data w.r.t mixture model



The process is iterated until convergence, ie when the likelihood of the data does not change (much)

Limitations

- ★ Hill-climbing \Rightarrow depends on initial parameters
- ★ Potential slow convergence, depending on the distributions
- ★ Hill-climbing \Rightarrow sensitive to local maxima

c-component mixtures

Generalisation with :

$$\Delta_{i1}, \Delta_{i2}, \dots, \Delta_{ik}, \dots, \Delta_{ic}$$

True (=1) if data x_i is generated by component ϕ_k

⇒ Responsibility γ_{ik} : expectation of Δ_{ik} over all the components

$\{\pi_k, \mu_k, \Sigma_k\}_{k=1,\dots,c}$ unknown parameters (before: $\pi_1 = \pi$, $\pi_2 = 1 - \pi$)

EM algorithm, c components

1. Initial $\theta^0 = \{\pi_k^0, \mu_k^0, \Sigma_k^0\}_{k=1, \dots, c}$
 - In general $\pi_k = 1/c$, μ_k is chosen at random and $\Sigma_k = \mathbf{Id}$
 - Alternative : use k-means as initialisation
2. **E-step** : compute responsibilities for every data $i = 1, \dots, N$ and every component $k = 1, \dots, c$

$$\gamma_{ik} = \frac{\pi_k \phi(x_i, \theta_k)}{\sum_{j=1}^c \pi_j \phi(x_i, \theta_j)}$$

3. **M-step** Estimations of mixture parameters

$$\mu_k = \frac{\sum_{i=1}^N \gamma_{ik} x_i}{\sum_i \gamma_{ik}}; \quad \Sigma_k = \frac{X_k \Gamma_k X_k^T}{\text{Tr}(\Gamma_k)}; \quad \pi_k = \frac{\sum_i \gamma_{ik}}{N}$$

with $\Gamma_k = \text{diag}[\gamma_{1k}, \dots, \gamma_{Nk}]$, X_k centered on μ_k

4. Iterate 2. and 3. until convergence

Modeling

- ★ The *a priori* parametrisation of the mixture changes the convergence
- ★ Parameters
 1. c : number of components
 2. Σ_k : the shape of covariance matrices (diagonal, full, parameterised)
- ★ Too flexible or too rigid models mean wrong or no convergence...
- ★ Number of variables : $p \times p \times c + 2 \times c$: if N low, p large and c large \rightarrow over-parameterised

Shape of the covariance matrix

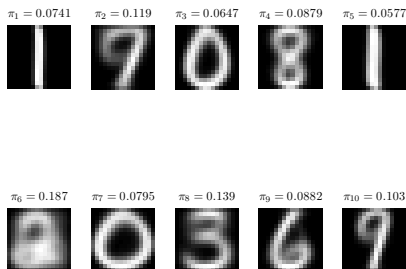
Over-parameterised problem

- ★ $\Sigma \in \mathbb{R}^{p \times p}$
 - ★ Eg: character recognition $\mathbf{x}_i \in \mathbb{R}^{256}$
- ⇒ Needs to estimate $256^2 \times c$ parameters for the covariance (given about 7000 data points)!

Matrix parameterisation

- ★ Spherical models $\Sigma = \sigma * \mathbf{Id}$, 1 parameter
- ★ Diagonal models $\Sigma = \text{diag}[\sigma_1, \dots, \sigma_p]$, p parameters
- ★ Full models $\Sigma \in \mathbb{R}^{p \times p}$, p^2 parameters

Character recognition



More complex models → no convergence since p is too large

Pre-processing : using PCA to reduce the dimension

Recall : 50 principal components reconstruct 90% of the signal

EM within the space of the 50 first PC



Pre-processing : using PCA to reduce the dimension

Recall : 50 principal components reconstruct 90% of the signal

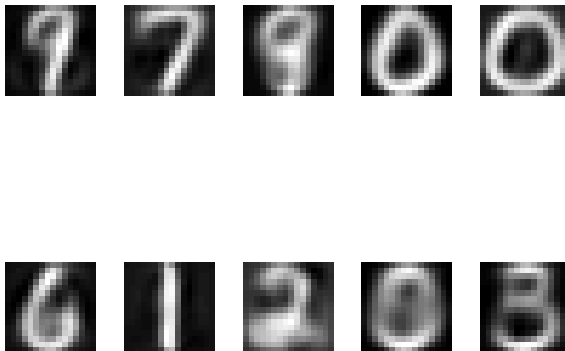
EM within the space of the 10 first PC



Pre-processing : using PCA to reduce the dimension

Recall : 50 principal components reconstruct 90% of the signal

EM within the space of the 2 first PC



Number of components

Parcimony

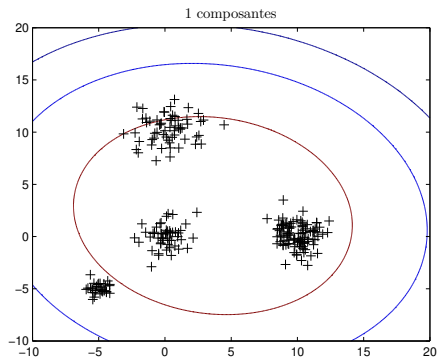
- ★ The larger c , the less points may be assigned to every component (in average)
- ★ Search for **parcimonious** models, ie small number of parameters to estimate

Bayesian Information Criterion (BIC)

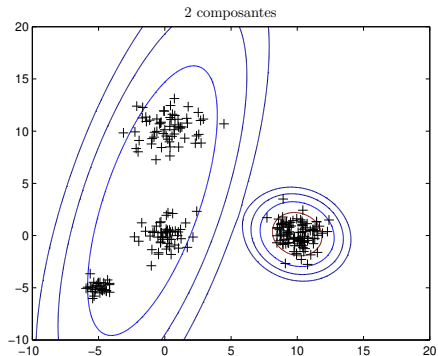
- ★ The larger c is, the better the estimate of l
- ★ Trade likelihood against complexity

$$BIC = 2 * l(\theta) - |\theta|. \log(N)$$

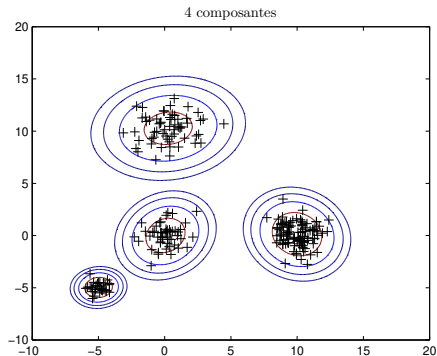
Example: 4-component mixture



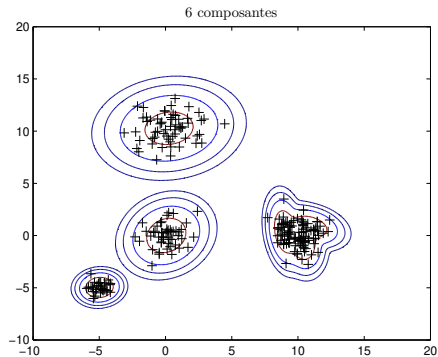
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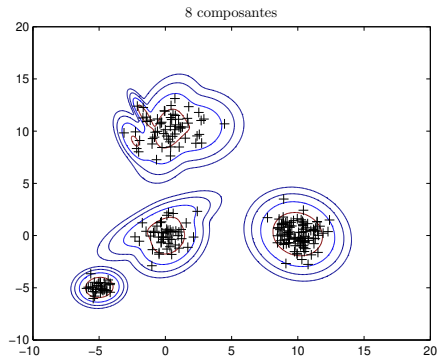
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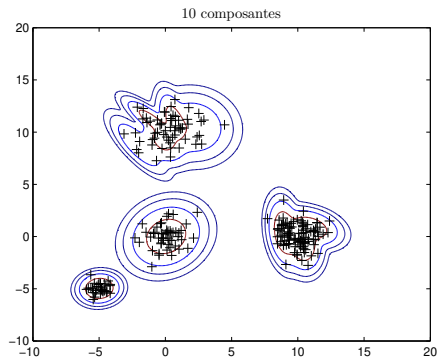
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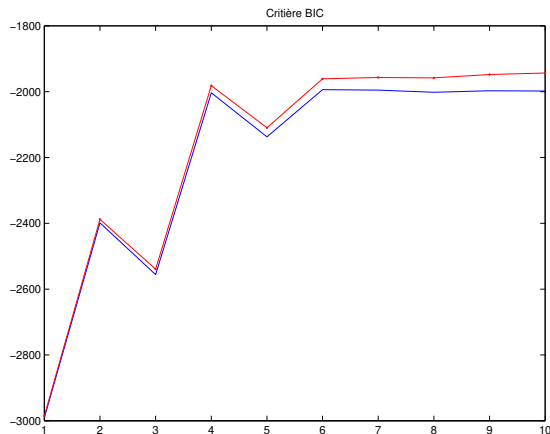
Example: 4-component mixture



Example: 4-component mixture



Example (cont'd)



- ★ Need to test all models
- ★ Depends on convergence
- ★ *Fine tuning* by hand!

Conclusions

Gaussian mixtures

- ★ The Gaussian mixture model generalises the underlying data hypothesis made by PCA and LDA
- ★ Explicit density modeling and estimation
- ★ Also **classification** (unsupervised): if components are classes, data point i is associated to class k for which $p(k|x_i) \approx \pi_k \phi_k(x_i)$ is maximised amongst all classes

Conclusions

EM algorithm

- ★ Iterative algorithm to maximise the (log-)likelihood
- ★ Principle used in many other scenarios
- ★ Based on the definition of hidden (**latent**) variables
- ★ Probabilistic Latent Semantic Analysis (pLSA) → EM where the hidden variables are the **latent concepts**