

Unsupervised Classification

A second partitioning-based algorithm: *Gaussian Mixture Models (GMM)*

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INSA Toulouse, Applied Mathematics, 4th year

1. Finite Mixture Models

- 1.1 General Principle
- 1.2 Step 1: Modeling and Choice of Distributions
- 1.3 Step 2: Parameters Estimation
- 1.4 Step 3: Maximum a Posteriori & Classification
- 1.5 Step 4: Model Selection Criteria

2. Gaussian Mixture Models

- 2.1 Multivariate Gaussian Mixtures

Finite Mixture Models

1.1 General Principle

1.2 Step 1: Modeling and Choice of Distributions

1.3 Step 2: Parameters Estimation

1.4 Step 3: Maximum a Posteriori & Classification

1.5 Step 4: Model Selection Criteria

Introduction

We observe n individuals described by p variables: $x_i = (x_{i1}, x_{i2}, \dots, x_{ip}) \in \mathcal{X}$

$$X = \begin{pmatrix} x_{11} & x_{12} & \dots & x_{1p} \\ x_{21} & x_{22} & \dots & x_{2p} \\ \vdots & \vdots & \ddots & \vdots \\ x_{n1} & x_{n2} & \dots & x_{np} \end{pmatrix}$$

$$\mathcal{X} = \mathbb{R}^p, \{0, 1\}^p,]-\pi, \pi]^p, \mathbb{R}^q \times \{0, 1\}^{p-q}, \dots$$

- Initial measurements,
 - Transformed measurements,
 - Coordinates after dimension reduction.
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- **Assumption:** The data come from a population composed of several sub-populations.

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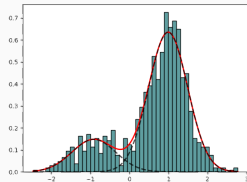
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- Initial measurements,
- Transformed measurements,
- Coordinates after dimension reduction.

- **Assumption:** The data come from a population composed of **several sub-populations**.

- **Modeling:**

- Each sub-population is modeled **independently of the others**,
 \rightsquigarrow Choice of a distribution law for each sub-population.
- Total population seen as a **mixture** of these sub-populations,
 \rightsquigarrow **Finite mixture model**.



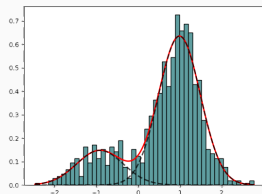
Finite Mixture Model

Mixture model of K components: *Given:*

- $(\alpha_k)_{k \in \llbracket 1, K \rrbracket}$ the *proportions* of the mixture

$$\forall k \in \llbracket 1, K \rrbracket, \alpha_k \in [0, 1] \quad \text{and} \quad \sum_{k=1}^K \alpha_k = 1;$$

- For all k , $f_k(\cdot; \omega_k)$ the *density* of the k -th sub-population, which (possibly) depends on a parameter ω_k ; *and*
- $\theta = (\alpha_k, \omega_k)_{k \in \llbracket 1, K \rrbracket}$ the whole *parameters* of the mixture model.



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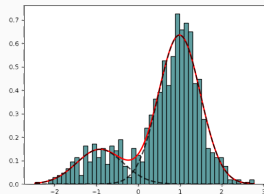
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We define:

$$q(\cdot; \theta) = \sum_{k=1}^K \alpha_k f_k(\cdot; \omega_k)$$



Mixture Models vs. Latent Variables

Mixture model: $q(\cdot; \theta) = \sum_{k=1}^K \alpha_k f_k(\cdot; \omega_k) \longleftrightarrow \begin{cases} \text{Proportions } \alpha_k \\ \text{Densities } f_k(\cdot; \omega_k) \end{cases}$

Question: *How to generate data according to such a model?* $(x_i)_{i \in \llbracket 1, n \rrbracket}$

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For all individual $i \in \llbracket 1, n \rrbracket$,

- (i) Let $\mathcal{P}_K = \{\mathcal{C}_1, \dots, \mathcal{C}_K\}$ be a partition of $\llbracket 1, n \rrbracket$ into K classes, such that the individual i belongs to \mathcal{C}_k with probability α_k : $\mathbb{P}(i \in \mathcal{C}_k; \alpha_k) = \alpha_k$;
- (ii) Then, x_i is generated according to the density $f_k(\cdot; \omega_k)$ if $i \in \mathcal{C}_k$.

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\leadsto **Latent variables** $(z_i)_{i \in \llbracket 1, n \rrbracket}$ to encode the classes.

For all individual $i \in \llbracket 1, n \rrbracket$,

$$\begin{cases} z_i \mid (\alpha_k)_{k \in \llbracket 1, K \rrbracket} \sim \sum_{k=1}^K \alpha_k \delta_k \\ x_i \mid z_i, \theta = (\alpha_k, \omega_k)_{k \in \llbracket 1, K \rrbracket} \sim f_{z_i}(\cdot; \omega_{z_i}) \end{cases}$$

Hierarchical Writing of Mixture Models

- **Mixture Models:** For all $i \in \llbracket 1, n \rrbracket$,
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- **Complete likelihood:** For all $x = (x_i)_{i \in \llbracket 1, n \rrbracket}$ and $z = (z_i)_{i \in \llbracket 1, n \rrbracket}$,

$$q(x, z; \theta) = \prod_{i=1}^n q(x_i, z_i; \theta) = \prod_{i=1}^n q(x_i \mid z_i; \theta) q(z_i; \theta) = \prod_{i=1}^n \alpha_{z_i} f_{z_i}(x_i; \omega_{z_i}).$$

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- **Posterior distribution:** For all $i \in \llbracket 1, n \rrbracket$,

$$\begin{aligned} q(x_i; \theta) &= \sum_{k=1}^K q(x_i, \{z_i = k\}; \theta) \\ &= \sum_{k=1}^K q(x_i \mid \{z_i = k\}; \theta) q(\{z_i = k\}; \theta) = \sum_{k=1}^K \alpha_k f_k(x_i; \omega_k) \end{aligned}$$

Steps to Define a Mixture Model

1. **Modeling and Choice of Distributions**
2. **Parameters Estimation**
3. **Maximum a Posteriori & Classification**
4. **Model Selection Criteria**

Steps to Define a Mixture Model

1. Modeling and Choice of Distributions

- Initial choice of the model,
- Selection of a suitable density family $f_k(\cdot; \omega_k)$,
- Choice to be made according to the problem/data studied,

↪ Collection of models: One model for each fixed number of classes $K \in \mathbb{N}^*$:

$$\mathcal{M}_K := \left\{ x \in \mathbb{R}^d \mapsto q_K(x; \theta) = \sum_{k=1}^K \alpha_k f_k(x; \omega_k) \right\}.$$

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- In each \mathcal{M}_K model, we identify the mixture that best fits the data: $q_K(\cdot; \hat{\theta})$.

↪ Parameter estimation algorithm.

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- Maximum A Posteriori (MAP) rule to derive a classification of the data.

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4. Model Selection Criteria

- Choose the “best” mixture model among $q_2(\cdot; \hat{\theta}), q_3(\cdot; \hat{\theta}), \dots, q_{K_{\max}}(\cdot; \hat{\theta})$

↪ Model selection criterion to determine \hat{K} and thus choose $q_{\hat{K}}(\cdot; \hat{\theta})$.

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Step 1: Modeling and Choice of Distributions

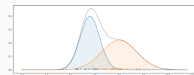
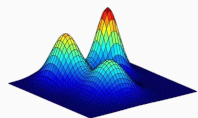
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 - A priori modeling, made **according to the problem/data** studied,
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- **Quantitative data:**
 - **Gaussian mixtures**,
 - Student mixtures, *etc.*
- **Qualitative data:**
 - Mixtures of multinomials, *etc.*
- **Counting data:**
 - Poisson mixtures,
 - Negative binomials, *etc.*
- **Compositional data:**
 - Dirichlet mixtures, *etc.*

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Step 2: Parameters Estimation – Maximum Likelihood Estimation

Assume a fixed number of classes $K \in \mathbb{N}^*$.

Shortcut: Let us note q for q_K .

- **Aim:** Maximize in $\theta = (\alpha_k, \omega_k)_{k \in \llbracket 1, K \rrbracket}$ the log-likelihood ℓ of the model.

(i) Recall that: $\forall i \in \llbracket 1, n \rrbracket, q(x_i; \theta) = \sum_{k=1}^K \alpha_k f_k(x_i; \omega_k)$

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\leadsto *Optimization algorithm to approximate $\hat{\theta}_{MLE}$.*

\implies **Expectation-Maximization (EM) algorithm:**

A core tool for estimation in latent variable models.

Step 2: Parameters Estimation – The EM Algorithm [Dempster et al., 1977]

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- Aim:** Given a latent variables model, find the MLE: $\hat{\theta}_{MLE} \in \operatorname{argmax}_{\theta \in \Theta} q(x; \theta)$

Observations: $X \mid x = (x_i)_{i \in \llbracket 1, n \rrbracket} \in \mathcal{X}$.

Latent variables: $Z \mid z = (z_i)_{i \in \llbracket 1, n \rrbracket} \in \mathcal{Z}$,

Here $z_i \in \llbracket 1, K \rrbracket$ denotes the class of the i -th individual.

Parameter: $\theta \in \Theta$, where Θ set of admissible parameters,

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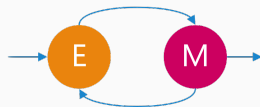
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- Principle:** Alternation of an *Expectation step* and a *Maximization step* until convergence.



Denote $\theta^{(t)}$ the current value of the θ parameter.

Step 2: Parameters Estimation – The EM Algorithm [Dempster et al., 1977]

E-step: Compute the conditional expected log-likelihood

$$\begin{aligned} Q(\theta|\theta^{(t)}) &= \int_{\mathcal{Z}} \log q(x, z; \theta) q(z|x; \theta^{(t)}) dz \\ &= \mathbb{E}_{Z \sim q(\cdot|x; \theta^{(t)})} [\log q(x, Z; \theta)] = \mathbb{E} [\log q(x, Z; \theta) | x; \theta^{(t)}] . \end{aligned}$$

M-step: Maximize $Q(\cdot|\theta^{(t)})$ in the feasible set Θ : $\theta^{(t+1)} \in \operatorname{argmax}_{\theta \in \Theta} Q(\theta|\theta^{(t)})$

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$$Q(\theta|\theta^{(t)}) = \sum_{i=1}^n \sum_{k=1}^K [\log(\alpha_k) + \log(f_k(x_i; \omega_k))] \tau_{ik}^{(t)}$$

where
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$$\begin{cases} \forall k \in \llbracket 1, K \rrbracket, & \alpha_k^{(t+1)} = \frac{1}{n} \sum_{i=1}^n \tau_{ik}^{(t)} \\ (\omega_k^{(t+1)})_{k \in \llbracket 1, K \rrbracket} \in & \operatorname{argmax}_{\omega = (\omega_k) \in \Omega} \sum_{i=1}^n \sum_{k=1}^K \tau_{ik}^{(t)} \log(f_k(x_i; \omega_k)) \end{cases}$$

The EM algorithm

E-step: Conditional expected log-likelihood

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Convergence of the EM algorithm

Suppose that:

- The f_k laws belong to the exponential family, and are sufficiently regular,
 - We can conduct the M-step at each iteration.
1. At every iteration of the EM algorithm, the log likelihood increases.
 2. The EM algorithm converges, but not necessarily to the global maximum of the log-likelihood, nor necessarily in finite time.

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In practice:

- Easy to implement,
- Sometimes slow to converge (especially when components are very mixed),
- Sensitive to the initialization, i.e. to the choice of $\theta^{(0)}$

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GEM: Generalized EM Algorithm [Delyon et al., 1999, Lange, 1995]

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2. Limitations concerning the **M-step**.
 - GEM:** Generalized EM Algorithm [Delyon et al., 1999, Lange, 1995]
3. Limitations concerning the **E-step**.
 - SEM:** Stochastic EM Algorithm [Celeux et al., 1996]
 - MCEM:** Monte-Carlo EM Algorithm [Wei and Tanner, 1990]
 - SAEM:** Stochastic-Approximation EM Algorithm [Delyon et al., 1999]

Variants of the EM Algorithm

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2. Limitations concerning the **M-step**.

GEM: Generalized EM Algorithm [Delyon et al., 1999, Lange, 1995]

3. Limitations concerning the **E-step**.

SEM: Stochastic EM Algorithm [Celeux et al., 1996]

MCEM: Monte-Carlo EM Algorithm [Wei and Tanner, 1990]

SAEM: Stochastic-Approximation EM Algorithm [Delyon et al., 1999]

SEM – Stochastic EM

S-step: Draw an unobserved sample

$$z^{(t)} \sim q(\cdot | x; \theta^{(t)})$$

“E”-step: Estim. of $Q(\cdot | \theta^{(t)})$

$$Q_t(\theta) = \log q(x, z^{(t)}; \theta)$$

M-step: Maximize Q_{t+1} :

$$\theta^{(t+1)} \in \operatorname{argmax}_{\theta \in \Theta} Q_{t+1}(\theta)$$

MCEM – Monte Carlo EM

S-step: Draw m samples

$$z_j^{(t)} \sim q(\cdot | x; \theta^{(t)})$$

“E”-step: Monte-Carlo estim.

$$Q_t(\theta) = \frac{1}{m} \sum_{j=1}^m \log q(x, z_j^{(t)}; \theta)$$

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SAEM – Stochastic Approx.

S-step: Draw a sample
 $z^{(t)} \sim q(\cdot | x; \theta^{(t)})$

SA-step: Update $Q_t(\theta)$ as
 $Q_{t+1}(\theta) = Q_t(\theta) + \gamma_t (\log q(x, z^{(t)}; \theta) - Q_t(\theta))$

M-step: Maximize Q_{t+1} :
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- EM type algorithms are sensitive to initialization.
- Search/Run/Select Strategy:
 - Choice of M initial positions,
 - Some iterations of the algorithm for each position,
 - Selection of the position with the highest likelihood.
- Initialize on the output of a k -means.
- Stochastic variants of the EM.
- More complicated strategies, *etc.*

Finite Mixture Models

- 1.1 General Principle
- 1.2 Step 1: Modeling and Choice of Distributions
- 1.3 Step 2: Parameters Estimation
- 1.4 **Step 3: Maximum a Posteriori & Classification**
- 1.5 Step 4: Model Selection Criteria

Step 3: Maximum a Posteriori & Classification

- **Principle:** Each individual is assigned to the class to which it has the **highest probability of belonging**, given the estimated parameter $\hat{\theta}_{MLE}$.

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Hence:

$$\tau_{ik}(\theta) = \frac{\alpha_k f_k(x_i; \omega_k)}{\sum_{\ell=1}^K \alpha_\ell f_\ell(x_i; \omega_\ell)}$$

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Hence:
$$\tau_{ik}(\theta) = \frac{\alpha_k f_k(x_i; \omega_k)}{\sum_{\ell=1}^K \alpha_\ell f_\ell(x_i; \omega_\ell)}$$

- **Maximum a Posteriori:** Let the estimate $\hat{\theta}_{MLE}$ of the parameter.

$$i \in \mathcal{C}_k \quad \text{iff} \quad \forall \ell \neq k, \quad \tau_{ik}(\hat{\theta}_{MLE}) > \tau_{i\ell}(\hat{\theta}_{MLE})$$

Example: One-dimensional Gaussian Mixture Model

Images from Victor Lavrenko.



Likelihood: $\theta = (\alpha, \mu_1, \mu_2, \sigma_1^2, \sigma_2^2)$

$$\begin{cases} q(x; \theta) = \alpha \phi(x; \mu_1, \sigma_1^2) + (1 - \alpha) \phi(x; \mu_2, \sigma_2^2) \\ \phi(x; \mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(x - \mu)^2}{2\sigma^2}\right) \end{cases}$$

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E-step: $Q(\theta|\theta^{(t)}) \longleftrightarrow \begin{cases} \tau_{i1}^{(t)} \\ \tau_{i2}^{(t)} = 1 - \tau_{i1}^{(t)} \end{cases}$

$$\tau_{i1}^{(t)} = \frac{\alpha^{(t)} \phi(x; \mu_1^{(t)}, \sigma_1^{2(t)})}{\alpha^{(t)} \phi(x; \mu_1^{(t)}, \sigma_1^{2(t)}) + (1 - \alpha^{(t)}) \phi(x; \mu_2^{(t)}, \sigma_2^{2(t)})}$$

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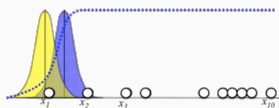
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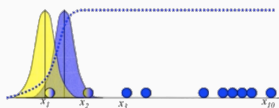
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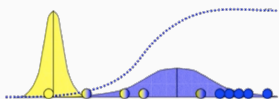
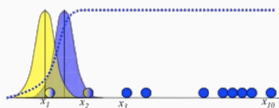
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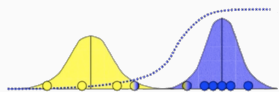
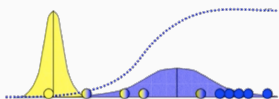
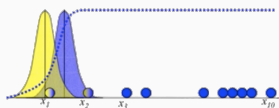
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CEM – Classification EM

E-step: Compute the $\tau_{ik}^{(t)}$

C-step: Determination of a partition of the data x by the MAP rule:

$$\hat{z}_i^{(t)} \in \operatorname{argmax}_{\ell \in \llbracket 1, K \rrbracket} \tau_{i\ell}^{(t)}$$

M-step: We update the parameters by replacing $\tau_{ik}^{(t)}$ by $\hat{z}_i^{(t)}$

- CEM converges in a **finite number of iterations**, contrary to EM,
- CEM produces biased estimators of the mixture parameters,
- We can prove that the k -means algorithm is a Gaussian mixture model with constant variance (ellipse \equiv circle), estimated by CEM.

Finite Mixture Models

- 1.1 General Principle
- 1.2 Step 1: Modeling and Choice of Distributions
- 1.3 Step 2: Parameters Estimation
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Step 4: Model Selection Criteria

- **Aim:** Determine the optimal number of classes K ,

$$\mathcal{M}_K = \left\{ x \in \mathbb{R}^d \mapsto q_K(x; \theta) = \sum_{k=1}^K \alpha_k f_k(x; \omega_k) \right\}.$$

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- **Probabilistic model**

↪ We can use the criteria from the [information theory](#):

$$\hat{K} = \underset{K}{\operatorname{argmin}} \operatorname{CRIT}(K) = \underset{K}{\operatorname{argmin}} \{-\log q_K(x; \theta) + \operatorname{pen}(K)\}.$$

AIC: Akaike Information Criterion [Akaike et al., 1973]

BIC: Bayesian Information Criterion [Schwarz, 1978]

ICL: Integrated Completed Likelihood [Biernacki et al., 2000]

Step 4: Model Selection Criteria

$$\hat{K} = \underset{K}{\operatorname{argmin}} \operatorname{CRIT}(K) = \underset{K}{\operatorname{argmin}} \{-\log q_K(x; \hat{\theta}_{MLE}) + \operatorname{pen}(K)\}.$$

AIC

$$\operatorname{AIC}(K) = -\log q_K(x; \hat{\theta}) + \nu_k$$

- Achieves a **bias-variance trade-off**
- Asymptotically, the AIC retains the model minimizing the *mean* Kullback deviation with the true unknown law
- In the context of finite mixture models, AIC tends to **under-penalize**

BIC

$$\operatorname{BIC}(K) = -\log q_K(x; \hat{\theta}) + \frac{\nu_k}{2} \log(n)$$

- Asymptotically, BIC selects the model minimizing the Kullback deviation from the true law.
 \rightsquigarrow The BIC is **convergent** if the true model is in the list of models.

ICL

$$\operatorname{ICL}(K) = -\log q_K(x, \hat{z}; \hat{\theta}) + \frac{\nu_k}{2} \log(n)$$

- Where \hat{z} is the MAP of $\hat{\theta}_{MLE}$

where ν_K is the number of free parameters of the mixtures \mathcal{M}_K .

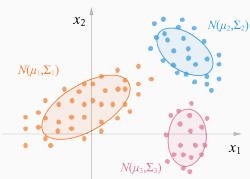
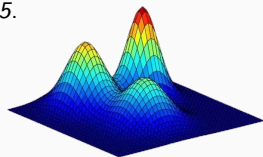
Gaussian Mixture Models

2.1 Multivariate Gaussian Mixtures

Multivariate Gaussian Mixtures

- Quantitative data: $x_i \in \mathbb{R}^d$, Generalization of slide 15.

- Likelihood: $\theta = (\alpha_k, \mu_k, \Sigma_k)_{k \in [1, K]}$



$$\left\{ \begin{array}{l} q_K(x; \theta) = \sum_{k=1}^K \alpha_k \phi(x; \mu_k, \Sigma_k) . \\ \phi(x; \mu, \Sigma) = \frac{1}{\sqrt{(2\pi)^d |\Sigma|}} \exp\left(-\frac{1}{2}(x - \mu)^\top \Sigma^{-1}(x - \mu)\right) . \end{array} \right.$$

- Gaussian Mixture Model:

$$\left\{ \begin{array}{l} \Theta_K = \left\{ (\alpha_k, \mu_k, \Sigma_k)_{k \in [1, K]} \in ([0, 1] \times \mathbb{R}^d \times \mathcal{S}_d \mathbb{R})^K \mid \sum_{k=1}^K \alpha_k = 1 \right\} \\ \mathcal{M}_K = \{ \theta \in \Theta_K \mid x \in \mathbb{R}^d \mapsto q_K(x; \theta) \} \end{array} \right.$$

- Estimation through the EM algorithm (See tutorials)

Estimation under Constraints

- Without constraint, $\dim(\Theta_K) = (K - 1) + Kd + K \frac{d(d + 1)}{2}$
 - ↪ In large dimensions, this can lead to an **over-parameterized** model.
 - ↪ Constraints on the **type of covariance** matrix.

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- Forms of Gaussian mixtures:**

Eigenvalue decomposition of covariance matrices:

$$\Sigma_k = L_k D_k A_k D_k^\top$$

- Volume:** $L_k = |\Sigma_k|^{1/d}$, *constant, or not.*
- Orientation:** D_k matrix of eigenvectors of Σ_k , *constant, or not.*
- Form:** A_k diagonal matrix of normalized eigenvectors of Σ_k , *spherical, diagonal or full.*

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- ↪ **14** type of covariance matrix:
- Spherical:** Equal volume, or not (2),
 - Diagonal:** Equal volume, or not ; Equal shape, or not (4),
 - Full:** Equal volume, or not ; Equal shape, or not ; Equal orientation, or not (8).

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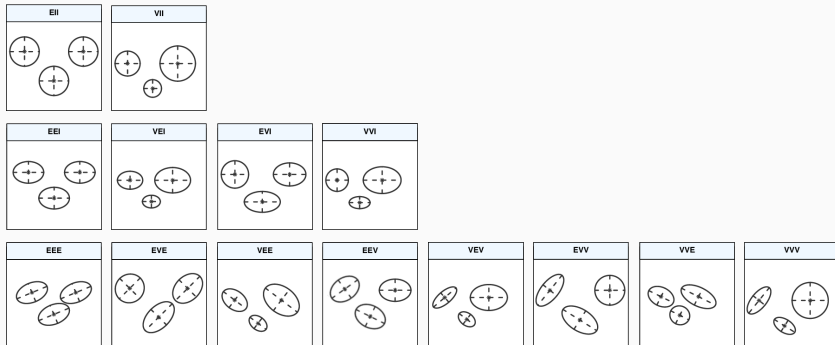
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Proportions α_k assumed equal or free.

↪ **28** possible forms of Gaussian mixtures.

Constraints on the Covariance Matrices

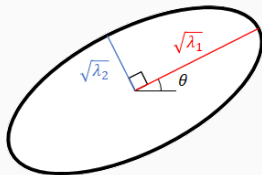
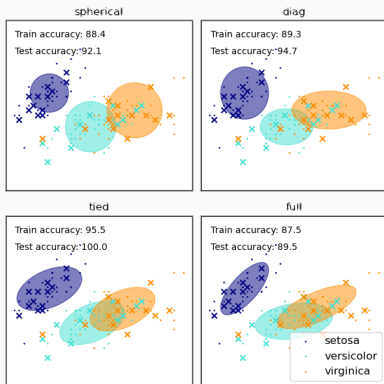
- Nomenclature in R: `modelName`s = `c("EEE", "VEE", "EVV", "VVV")`.
(Package `mclust`, page 38 for a description of all types.)



- In Python: “full”, “tied”, “diag”, “spherical”.
(Function `sklearn.mixture.GaussianMixture`)

Covariance Matrices vs. Rotation Matrices in \mathbb{R}^2

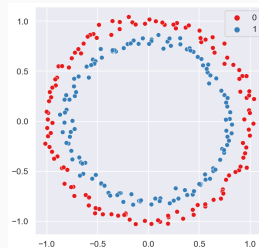
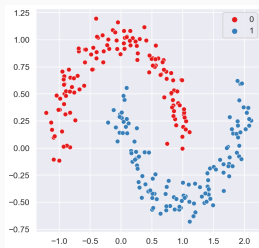
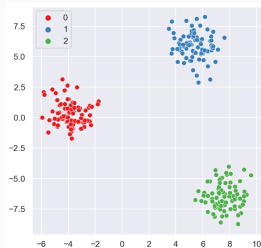
- D : Rotation matrix defined by an angle α : $D = \begin{pmatrix} \cos \alpha & \sin \alpha \\ -\sin \alpha & \cos \alpha \end{pmatrix}$,
- A : Diagonal matrix of diagonal terms b and $1/b$: $A = \begin{pmatrix} b & 0 \\ 0 & \frac{1}{b} \end{pmatrix}$,
- Ellipse of equidensity: $L = \lambda = \text{volume}$.



$$\begin{cases} \lambda_1 = \lambda b \\ \lambda_2 = \frac{\lambda}{b} \end{cases}$$

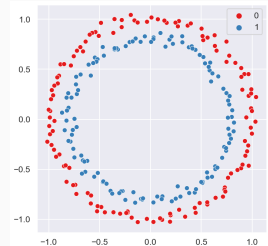
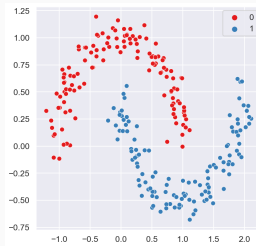
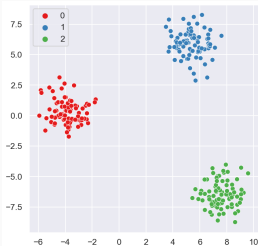
Strengths and Weaknesses

- Pros:**
- Give probabilistic cluster assignments,
 - Have probabilistic interpretation,
 - Can handle clusters with varying sizes, variance, *etc.*



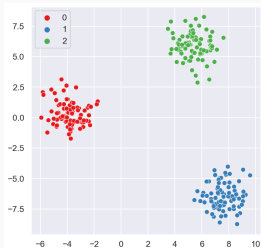
Strengths and Weaknesses

- Pros:**
- Give probabilistic cluster assignments,
 - Have probabilistic interpretation,
 - Can handle clusters with varying sizes, variance, *etc.*
- Cons:**
- Initialization matters,
 - Choose appropriate distributions,
 - Overfitting issues.

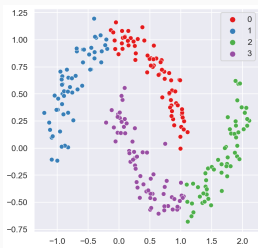


Strengths and Weaknesses

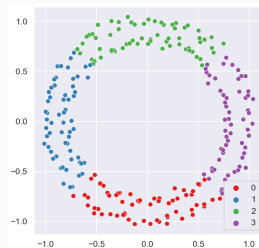
- Pros:**
- Give probabilistic cluster assignments,
 - Have probabilistic interpretation,
 - Can handle clusters with varying sizes, variance, *etc.*
- Cons:**
- Initialization matters,
 - Choose appropriate distributions,
 - Overfitting issues.



Tied, 3 components



Full, 4 components



Diagonal, 4 components

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