Clustering

Course of Machine Learning
Master Degree in Computer Science
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Partitional clustering

Problem
Given a dataset $\mathbf{X} = (\mathbf{x}_1, \ldots, \mathbf{x}_n)$, with $\mathbf{x}_i \in \mathbf{R}^d (i = 1, \ldots, n)$.

We wish to derive a set of clusters *clusters* (i.e. a partition of $\mathbf{X}$ into subsets of "near" elements). Clusters are represented by their *prototypes* $(\mathbf{m}_1, \ldots, \mathbf{m}_k)$, with $\mathbf{m}_j \in \mathbf{R}^d, j = 1, \ldots, k$.

Rappresentation of a clustering

1. Cluster prototypes $(\mathbf{m}_1, \ldots, \mathbf{m}_k)$, with $\mathbf{m}_j \in \mathbf{R}^d (j = 1, \ldots, k)$
2. Element assignment to clusters: for each $\mathbf{x}_i$, k binary flags $r_{ij} \in \{0, 1\}$, $j = 1, \ldots, k$. If $\mathbf{x}_i$ is assigned the t-th cluster, then $r_{it} = 1$ and $r_{ij} = 0$ for $j \neq t$

Clustering types

Partitional clustering
Given a set of items (points) $\mathbf{X} = \{\mathbf{x}_1, \ldots, \mathbf{x}_n\}$, we wish to partition $\mathbf{X}$ by assigning each element to one out of k clusters $C_1, \ldots, C_k$ in such a way to maximize (or minimize) a given cost J. The number k of clusters could be given or should have to be computed.

Hierarchical clustering
Given a set of items (points) $\mathbf{X} = \{\mathbf{x}_1, \ldots, \mathbf{x}_n\}$, we wish to derive a set of nested partitions of $\mathbf{X}$, from the partition composed by all singletons (one cluster for each node) to the one composed by a single item (the whole set).

Partitional clustering

Brute force methods
Check all partitions of a set of n elements into k subsets, selecting the one with minimum J. The number $P(n, k)$ of such partitions can be recursively defined as follows:

$$\begin{aligned} P(n+1, k) &= P(n, k-1) + kP(n, k) \\ P(n, 1) &= 1 \\ P(n, n) &= 1 \end{aligned}$$

It is possible to prove that this results in the following closed form characterization:

$$P(n, k) = \frac{1}{k!} \sum_{j=0}^{k} (-1)^{k-j} \binom{k}{j} j^n$$

This is the *Stirling number* of the second type which is known to be at least $\frac{1}{2}(k^2+k+2)k^{n-k-1}$ for $n \geq 2$, $1 \leq k \leq n-1$.

Clustering cost

Sum of squares

Let us define the cost a clustering as follows:

$$J(R,M) = \sum_{i=1}^{k}\sum_{j=1}^{n} r_{ij}||\mathbf{x}_j - \mathbf{m}_i||^2 = \sum_{i=1}^{k}\sum_{j=1}^{n} r_{ij}(\mathbf{x}_j - \mathbf{m}_i)^T(\mathbf{x}_j - \mathbf{m}_i)$$

where

- $R_{ij} = r_{ij}$, where $r_{is} = 1$ and $r_{ij} = 0$ for $j \neq s$ if x_i is assigned to cluster C_s
- $M_i = \mathbf{m}_i$, $i = 1, \ldots, k$ is the prototype (centroid) of cluster C_i,

$$\mathbf{m}_i = \frac{1}{n_i}\sum_{j=1}^{n} r_{ij}\mathbf{x}_j$$

k-means clustering

Dataset $\mathbf{X} = (x_1, \ldots, x_n)$, $x_i \in \mathbf{R}^d$: we wish to derive k clusters with prototypes $\mathbf{m}_1, \ldots, \mathbf{m}_k$

Assignment of elements to cluster: for each x_i, k binary flags r_{ij} ($j = 1, \ldots, k$)

- if x_i is assigned to cluster s, then $r_{is} = 1$, and $r_{ij} = 0$ for $j \neq k$

Cost: sum of the distances of each point from the prototype of the corresponding cluster

$$J(R,M) = \sum_{i=1}^{n}\sum_{j=1}^{k} r_{ij}\,||x_i - \mathbf{m}_j||^2$$

Objective: finding r_{ij} and $\mathbf{m}_j$ ($i = 1, \ldots, n$, $j = 1, \ldots, k$) to minimize $J(R,M)$

Algorithm

1. Given a set of prototypes $\mathbf{m}_{ij}$, minimize wrt r_{ij} (assigning elements to clusters).

   For each x_i, minimize $\sum_{j=1}^{k} r_{ij}\,||x_i - \mathbf{m}_j||^2$.

   The minimum is obtained for $r_{ik} = 1$ (and $r_{ij} = 0$ for $j \neq k$), where $||x_i - \mathbf{m}_k||^2$ is the minimum distance. That is, each point is assigned to the cluster of the nearest prototype.

2. Given a set of assignments r_{ij}, minimize wrt $\mathbf{m}_{ij}$ (defining new cluster prototypes)

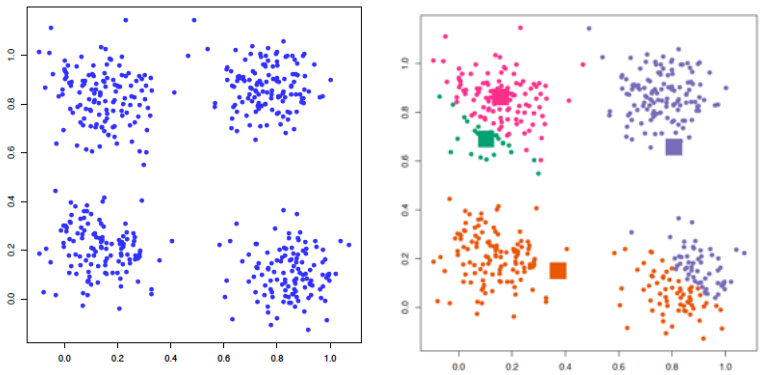
   For each$\mathbf{m}_k$, $J = \sum_{i=1}^{n}\sum_{j=1}^{k} r_{ij}\,||x_i - \mathbf{m}_j||^2$ is a quadratic function of $\mathbf{m}_k$. By setting its derivative to zero, the values of $\mathbf{m}_k$ providing its minimum are obatined

   $$\frac{\partial J}{\partial \mathbf{m}_k} = 2\sum_{i=1}^{n} r_{ik}(x_i - \mathbf{m}_k) = 0 \Longrightarrow \mathbf{m}_k = \frac{\sum_{i=1}^{n} r_{ik}x_i}{\sum_{i=1}^{n} r_{ik}}$$

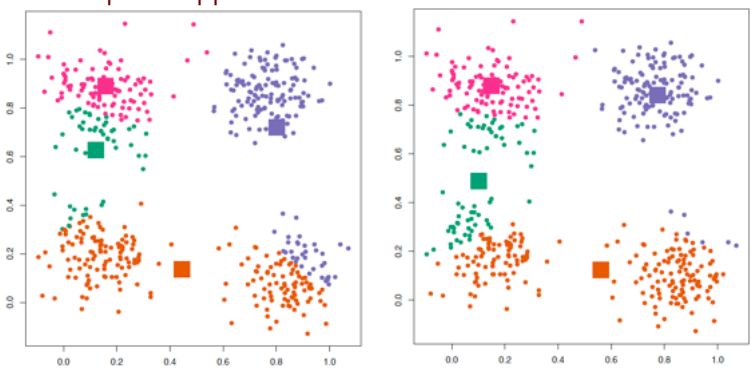
   That is, the new prototype is the mean of the elements assigned to the cluster

At each step, J does not increase. There is a convergence to a local minimum.

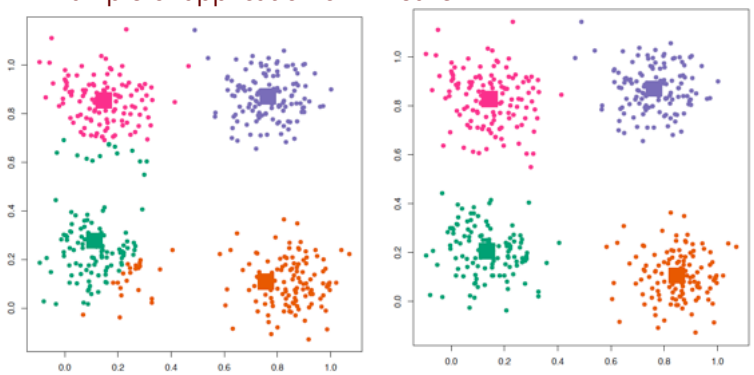
Example of application of k-means



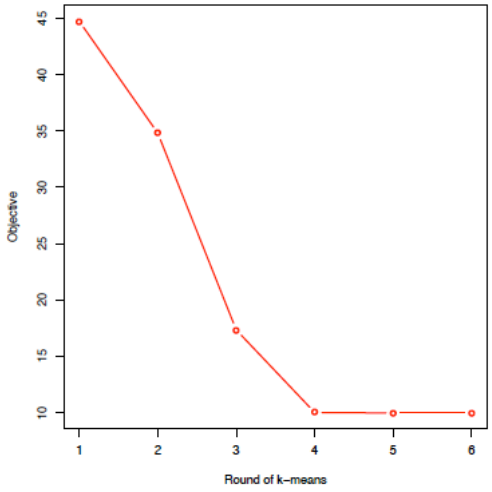
Example of application of k-means



Example of application of k-means



Example of application of k-means

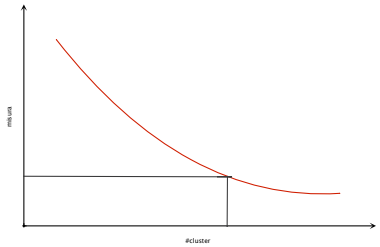


How to choose K

Cross validation

- Apply cross validation for different values of K, measuring the quality of the clustering obtained
- How to measure the quality of a clustering?
  1. mean distance of elements from the prototypes of their clusters
  2. log-likelihood of the elements wrt the resulting mixture model

Note
Measures improves as K increases (overfitting). A value such that further increases provide limited improvement should be found



How to choose K

Penalty
Use of penalty terms wrt number of parameters

- Akaike Information Criterion (AIC)

$$\text{AIC} = 2K - 2\ln L$$

- Bayesian Information Criterion (BIC)

$$\text{BIC} = K\ln n - 2\ln L$$

where L is the model likelihood

Hierarchical clustering

Aim
Derivation of a binary tree. Node: cluster; arc: inclusion.
The tree specifies a set of pairwise merge of clusters.

- Aggregation, starting from n singleton clusters
- Separation, starting from a single cluster of size n

Requirements
k-means requires:

- a number K of clusters
- an initizl assignment
- a distance function between elements

Hierarchical clustering requires:

- a similarity function between clusters

Hierarchical clustering by aggregation

Algorithm

- define n clusters (singleton)
- repeat
  - compute the matrix of distances between clusters
  - merge the pair of clusters which are "nearest"
- until "a single cluster has remained"

Hierarchical clustering by aggregation

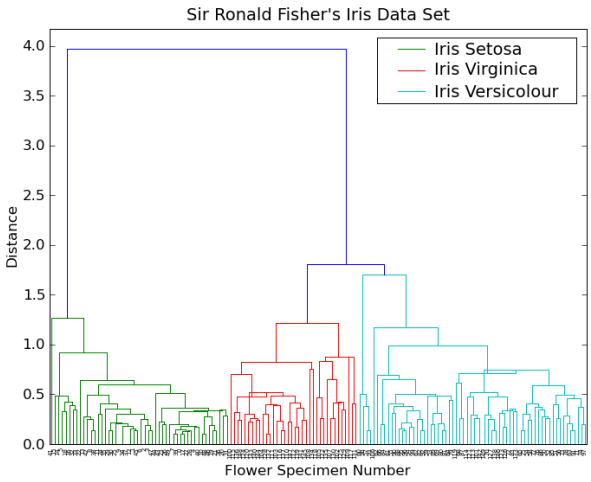
Properties

- Each tree level is a partition of elements
- The algorithm provides a sequence of clusterings
- The best clustering has to be found
- Monotonicity: similarity between paired clusters decreases

Dendrogram

- Tree of cluster pairings
- The height of the nodes is inversely proportional to the similarity of the paired clusters

Dendrogramma



Cluster similarity

Many measures. Most frequent ones:

- Similarity between nearest nodes (*Single linkage*)

$$d_{SL}(C_1, C_2) = \min_{\mathbf{x}_1 \in C_1, \mathbf{x}_2 \in C_2} d(x_i, x_j)$$

- Similarity between farthest nodes (*Complete linkage*)

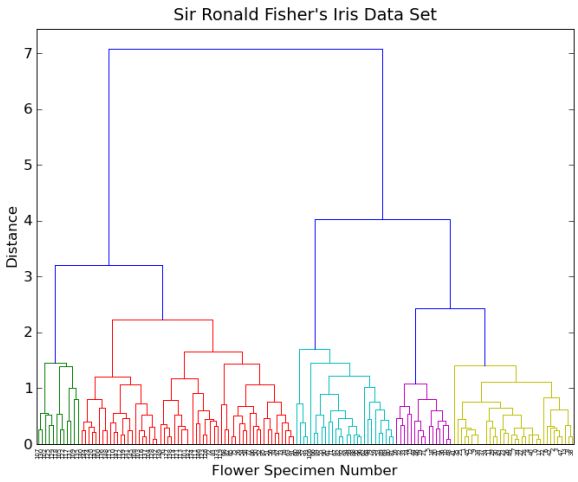
$$d_{CL}(C_1, C_2) = \max_{\mathbf{x}_1 \in C_1, \mathbf{x}_2 \in C_2} d(x_i, x_j)$$

- Mean similarity (*Group average*)

$$d_{GA}(C_1, C_2) = \frac{1}{|C_1| \cdot |C_2|} \sum_{\mathbf{x}_1 \in C_1} \sum_{\mathbf{x}_2 \in C_2} d(x_i, x_j)$$

Different measures provide different dendrograms

Dendrogram with complete linkage



Mixtures of distributions

Linear combinations of probability distributions $q(x|\theta)$

- Same type of distributions
- Differ by parameter values

$$p(x|\boldsymbol{\pi}, \boldsymbol{\theta}) = \sum_{k=1}^{K} \pi_k q(x|\theta_k)$$

where

$$\boldsymbol{\pi} = (\pi_1, \ldots, \pi_K) \qquad \boldsymbol{\theta} = (\theta_1, \ldots, \theta_K)$$

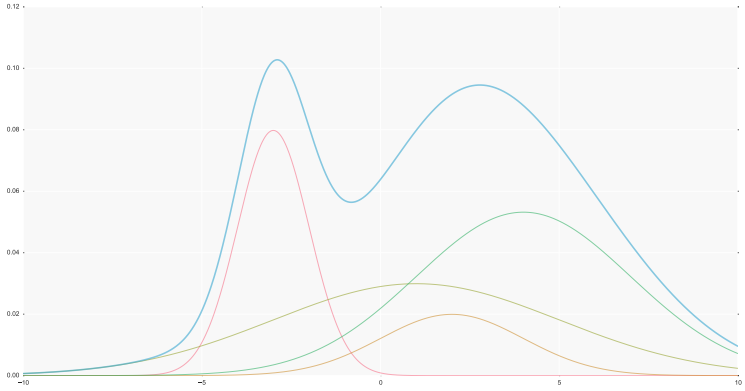
Mixing coefficients

$$0 \leq \pi_k \leq 1 \quad k = 1, \ldots, K \qquad \sum_{k=1}^{K} \pi_k = 1$$

Terms π_k have the properties of probability values

Mixtures of distributions

Provide extensive capabilities to model complex distributions. For example, almost all continuous distributions can be modeled by the linear combination of a suitable number of gaussians.



Mixture parameters estimation

Given a dataset $\mathbf{X} = (x_1, \ldots, x_n)$, the parameters $\boldsymbol{\pi}, \boldsymbol{\theta}$ of a mixture can be estimated by maximum likelihood.

$$L(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X}) = p(\mathbf{X}|\boldsymbol{\theta}, \boldsymbol{\pi}) = \prod_{i=1}^{n} p(x_i|\boldsymbol{\theta}, \boldsymbol{\pi}) = \prod_{i=1}^{n} \sum_{k=1}^{K} \pi_k q(x|\theta_k)$$

or maximum log-likelihood

$$l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X}) = \log p(\mathbf{X}|\boldsymbol{\theta}, \boldsymbol{\pi}) = \sum_{i=1}^{n} \log p(x_i|\boldsymbol{\theta}, \boldsymbol{\pi}) = \sum_{i=1}^{n} \log \left(\sum_{k=1}^{K} \pi_k q(x_i|\theta_k) \right)$$

Mixture parameters estimation

Let us derive the set of derivatives for $j = 1, \ldots, K$ and set them to 0

$$\frac{\partial l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X})}{\partial \theta_j} = \frac{\partial}{\partial \theta_j} \left[\sum_{i=1}^{n} \log \left(\sum_{k=1}^{K} \pi_k q(x_i|\theta_k) \right) \right] = 0$$

$$\frac{\partial l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X})}{\partial \pi_j} = \frac{\partial}{\partial \pi_j} \left[\sum_{i=1}^{n} \log \left(\sum_{k=1}^{K} \pi_k q(x_i|\theta_k) \right) \right] = 0$$

which itself results, for $k = 1, \ldots, K$, into

$$\pi_k = \frac{1}{n} \sum_{i=1}^{n} \gamma_k(x_i) \qquad \sum_{i=1}^{n} \gamma_k(x_i) \frac{\partial \log q(x_i|\theta_k)}{\partial \theta_k} = 0$$

where

$$\gamma_k(x) = \frac{\pi_k q(x|\theta_k)}{\sum_{j=1}^{K} \pi_j q(x|\theta_j)}$$

Mixture parameters estimation

The constraint $\sum_{i=1}^{K} \pi_i = 0$ can be taken into account by introducing a Lagrange multiplier λ and considering the Lagrangian

$$L(\boldsymbol{\theta}, \boldsymbol{\pi}, \lambda) = l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X}) + \lambda(1 - \sum_{i=1}^{K} \pi_i)$$

Setting the derivative wrt π_j to 0 turns out to be equivalent to

$$\lambda = \frac{\partial l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X})}{\partial \pi_j} = \frac{\partial}{\partial \pi_j}\left[\sum_{i=1}^{n} \log\left(\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)\right)\right] = \sum_{i=1}^{n} \frac{\partial}{\partial \pi_j}\left[\log\left(\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)\right)\right]$$
$$= \sum_{i=1}^{n} \frac{1}{\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)} \frac{\partial}{\partial \pi_j}\left(\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)\right)$$
$$= \sum_{i=1}^{n} \frac{1}{\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)} \sum_{k=1}^{K} \frac{\partial}{\partial \pi_j}\left(\pi_k q(x_i|\theta_k)\right)$$
$$= \sum_{i=1}^{n} \frac{q(x_i|\theta_j)}{\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)} = \sum_{i=1}^{n} \frac{\gamma_j(x_i)}{\pi_j} = \frac{1}{\pi_j} \sum_{i=1}^{n} \gamma_j(x_i)$$

Mixture parameters estimation

Setting the derivative wrt λ to 0

$$\frac{\partial}{\partial \lambda}\left(l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X}) + \lambda(1 - \sum_{i=1}^{K} \pi_i)\right) = 0$$

is equivalent to

$$\sum_{i=1}^{K} \pi_i = 1$$

Moreover, since, as shown above,

$$\pi_j = \frac{1}{\lambda} \sum_{i=1}^{n} \gamma_j(x_i)$$

it results

$$\sum_{j=1}^{K} \pi_j = \frac{1}{\lambda} \sum_{j=1}^{K} \sum_{i=1}^{n} \gamma_j(x_i) = 1$$

and

$$\lambda = \sum_{j=1}^{K} \sum_{i=1}^{n} \gamma_j(x_i) = \sum_{i=1}^{n} \sum_{j=1}^{K} \gamma_j(x_i) = \sum_{i=1}^{n} \sum_{j=1}^{K} \frac{\pi_j q(x_i|\theta_j)}{\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)} = \sum_{i=1}^{n} 1 = n$$

Mixture parameters estimation

Finally,

$$\begin{aligned}
\frac{\partial l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X})}{\partial \theta_j} &= \frac{\partial}{\partial \theta_j}\left[\sum_{i=1}^{n} \log\left(\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)\right)\right] = \sum_{i=1}^{n} \frac{\partial}{\partial \theta_j}\left[\log\left(\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)\right)\right] \\
&= \sum_{i=1}^{n} \frac{1}{\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)} \frac{\partial}{\partial \theta_j}\left(\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)\right) \\
&= \sum_{i=1}^{n} \frac{1}{\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)} \sum_{k=1}^{K} \frac{\partial}{\partial \theta_j}\left(\pi_k q(x_i|\theta_k)\right) \\
&= \sum_{i=1}^{n} \frac{\pi_j}{\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)} \frac{\partial}{\partial \theta_j} q(x_i|\theta_j) \\
&= \sum_{i=1}^{n} \frac{\pi_j q(x_i|\theta_j)}{\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)} \frac{1}{q(x_i|\theta_j)} \frac{\partial}{\partial \theta_j} q(x_i|\theta_j) \\
&= \sum_{i=1}^{n} \frac{\pi_j q(x_i|\theta_j)}{\sum_{k=1}^{K} \pi_k q(x_i|\theta_k)} \frac{\partial \log q(x_i|\theta_j)}{\partial \theta_j} = \sum_{i=1}^{n} \gamma_j(x_i) \frac{\partial \log q(x_i|\theta_j)}{\partial \theta_j} = 0
\end{aligned}$$

Mixture parameters estimation

Log likelihood maximization is intractable analytically: its solution cannot be given in closed form.

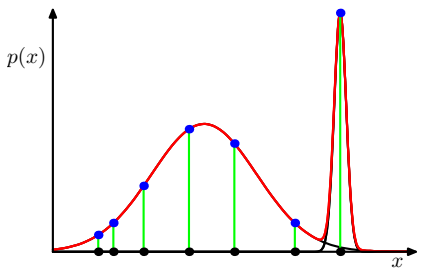
- $\boldsymbol{\pi}$ and $\boldsymbol{\theta}$ can be derived from $\gamma_k(x_i)$
- Also, $\gamma_k(x_i)$ can be derived from $\boldsymbol{\pi}$ e $\boldsymbol{\theta}$

Iterative techniques

- Given an estimation for $\boldsymbol{\pi}$ e $\boldsymbol{\theta}$...
- derive an estimation for $\gamma_k(x_i)$, from which ...
- derive a new estimation for $\boldsymbol{\pi}$ e $\boldsymbol{\theta}$, from which ...
- derive a new estimation for $\gamma_k(x_i)$ ...

Issues in ML for mixtures

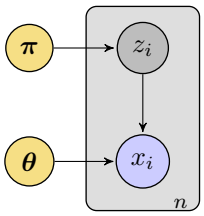
- Identifiability: for each solution (assignment of parameters to component distributions), there exist $K! - 1$ equivalent solutions
- Singularity: risk of severe overfitting. A mixture collapses to a single point.



Mixtures as generative processes

Graphical model representation of a mixture of distributions.

Latent variables



- Terms z_i are *latent* random variable with domain $z \in \{1, \dots, K\}$
- While x_i is observed, the value of z_i cannot be observed
- z_i denotes the component distribution $q(x|\theta)$ responsible for the generation of x_i

Mixtures as generative processes

Generation process

1. Starting from the distribution $\pi_1, \dots, \pi_K$, the component distribution to apply to sample the value of x_i is sampled: its index is given by z_i: hence z_i is dependent from $\boldsymbol{\pi}$
2. Let $z_i = k$: then, x_i is sampled from distribution $q(x|\theta_k)$. That is, x_i is dependent from both z_i and $\boldsymbol{\theta}$

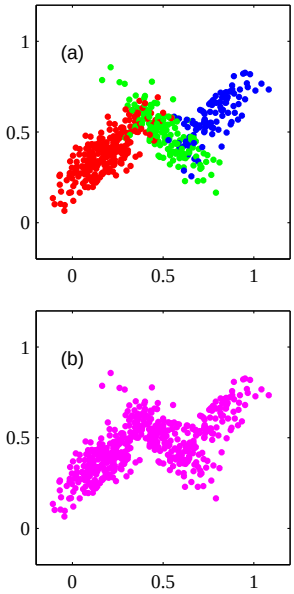
Latent variables coding

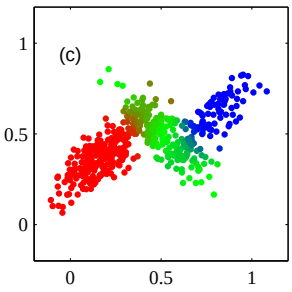
Indeed, z_i can be seen as components of a single latent K-dimensional variable $\boldsymbol{\zeta} = (\zeta_1, \dots, \zeta_K)$

1-to-K coding: K possible values $\zeta_i \in \{0, 1\}$, $\sum_{i=1}^{K} \zeta_i$.

Mixtures as generative processes

Example of generation of dataset from mixture of 3 gaussians





Distributions with latent variables

$$p(x|z=k,\boldsymbol{\theta},\boldsymbol{\pi}) = p(x|z=k,\boldsymbol{\theta}) = q(x|\theta_k)$$

Marginalizing wrt z,

$$\begin{aligned} p(x|\boldsymbol{\theta},\boldsymbol{\pi}) &= \sum_{k=1}^{K} p(x,z=k|\boldsymbol{\theta},\boldsymbol{\pi}) = \sum_{k=1}^{K} p(x|z=k,\boldsymbol{\theta})p(z=k|\boldsymbol{\pi}) \\ &= \sum_{k=1}^{K} q(x|\theta_k)p(z=k|\boldsymbol{\pi}) \end{aligned}$$

Since, by definition,

$$p(x|\boldsymbol{\theta},\boldsymbol{\pi}) = \sum_{k=1}^{K} \pi_k q(x_i|\theta_k)$$

it results

$$p(z=k|\boldsymbol{\theta},\boldsymbol{\pi}) = p(z=k|\boldsymbol{\pi}) = \pi_k$$

Responsibilities
An interpretation for $\gamma_k(x)$ can be derived as follows

$$\begin{aligned} \gamma_k(x) &= \frac{\pi_k q(x|\theta_k)}{\sum_{j=1}^{K} \pi_j q(x|\theta_j)} \\ &= \frac{p(z=k)p(x|z=k)}{\sum_{j=1}^{K} p(z=j)p(x|z=j)} = p(z=k|x) \end{aligned}$$

Mixing coefficients and responsibilities

- A mixing coefficient $\pi_k = p(z=k)$ can be seen as the prior (wrt to the observation of the point) probability that the next point is generated by sampling the k-th component distribution
- A responsibility $\gamma_k(x) = p(z=k|x)$ can be seen as the posterior (wrt to the observation of the point) probability that a point has been generated by sampling the k-th component distribution

Maximum likelihood

Data set

- Let $\mathbf{X} = (x_1, \ldots, x_n)$ be the set of values of observed variables and let $\mathbf{Z} = (z_1, \ldots, z_n)$ be the set of values of the latent variables. Then $(\mathbf{X}, \mathbf{Z})$ is the *complete dataset*: it includes the values of all variables in the model
- $\mathbf{X}$ is the *observed dataset* (incomplete). It only includes "real" data, that is observed data.

Indeed, $\mathbf{Z}$ is unknown. If values have been assigned to model parameters, the only possible knowledge about $\mathbf{Z}$ is given by the posterior distribution $p(\mathbf{Z}|\mathbf{X}, \boldsymbol{\theta}, \boldsymbol{\pi})$.

Inferring parameters for gaussian mixtures

- If we assume that the complete dataset $(\mathbf{X}, \mathbf{Z})$ is known (that is the observed points *together with their corresponding components*) a maximum likelihood estimation of $\boldsymbol{\pi}$ and $\boldsymbol{\theta}$ would be easy. In particular,
- For the mixing coefficients π_k it would result, as usual

  $$\pi_k = \frac{n_k}{n}$$

  where n_k is the number of elements of the set C_k such that $z = k$
- For component parameters $\theta_k = (\mu_k, \Sigma_k)$ the usual estimations for gaussians would provide

  $$\boldsymbol{\mu}_k = \frac{1}{n_k} \sum_{\mathbf{x} \in C_k} \mathbf{x}$$
  $$\boldsymbol{\Sigma}_k = \frac{1}{n_k} \sum_{\mathbf{x} \in C_k} (\mathbf{x} - \boldsymbol{\mu}_k)(\mathbf{x} - \boldsymbol{\mu}_k)^T$$

Log likelihood of complete dataset

The above results derive from the maximimization, wrt π_k, μ_k, Σ_k, $(k = 1, \ldots, K)$ of the log likelihood

$$\begin{aligned} l(\boldsymbol{\psi}|\mathbf{X}, \mathbf{Z}) = \log p(\mathbf{X}, \mathbf{Z}|\boldsymbol{\theta}, \boldsymbol{\pi}) &= \log \prod_{i=1}^{n} \prod_{k=1}^{K} \pi_k^{z_{ik}} \mathcal{N}(x_i|\mu_k, \Sigma_k)^{z_{ik}} \\ &= \sum_{i=1}^{n} \sum_{k=1}^{K} z_{ik} (\log \pi_k + \log \mathcal{N}(x_i|\mu_k, \Sigma_k)) \end{aligned}$$

Dealing with latent variables

Unfortunately, since $\mathbf{Z}$ is unknown, the log-likelihood of the complete dataset cannot be defined (the sets C_k are not known).

Our approach will be to consider for maximization, instead of the log-likelihood where each z_i is specified,

- its expectation wrt to the conditional distribution $p(\mathbf{Z}|\mathbf{X})$, that is

$$\begin{aligned} E_{p(\mathbf{Z}|\mathbf{X})}[l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X}, \mathbf{Z})] &= \sum_{i=1}^{n} \sum_{k=1}^{K} p(z_i = k|\mathbf{x}_i)(\log \pi_k + \log q(\mathbf{x}_i|\theta_k)) \\ &= \sum_{i=1}^{n} \sum_{k=1}^{K} \gamma_k(\mathbf{x}_i)(\log \pi_k + \log q(\mathbf{x}_i|\theta_k)) \end{aligned}$$

Observe that this expectation can be derived if $p(\mathbf{Z}|\mathbf{X})$ (that is the set of all values $\gamma_k(\mathbf{x}_i)$) is known.

Maximization of expected log-likelihood

The maximization of $E_{p(\mathbf{Z}|\mathbf{X})}[l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X}, \mathbf{Z})]$ wrt to π_k, μ_k, Σ_k results easily into

$$\pi_k = \frac{1}{n}\sum_{i=1}^{n}\gamma_k(\mathbf{x}_j)$$

$$\boldsymbol{\mu}_k = \frac{1}{n_k}\sum_{i=1}^{n}\gamma_k(\mathbf{x}_i)\mathbf{x}_i$$

$$\boldsymbol{\Sigma}_k = \frac{1}{n_k}\sum_{i=1}^{n}\gamma_j(\mathbf{x}_i)(\mathbf{x}_i - \boldsymbol{\mu}_k)(\mathbf{x}_i - \boldsymbol{\mu}_k)^T$$

this is named *M-step* (from "Maximization")

A new expectation

The computed values for the parameters result into new, different values for $\gamma_k(x_i) = p(z_i = k|x_i)$, and a different expectation $E_{p(\mathbf{Z}|\mathbf{X})}[l(\boldsymbol{\theta}, \boldsymbol{\pi}|\mathbf{X}, \mathbf{Z})]$.

$$p(z_i = k|x_i) = \frac{p(z_i = k)p(x_i|z_i = k)}{\sum_{j=1}^{K} p(z_i = j)p(x_i|z_i = j)} = \frac{\pi_k \mathcal{N}(x_i|\mu_k, \Sigma_k)}{\sum_{j=1}^{K} \pi_j \mathcal{N}(x_i|\mu_j, \Sigma_j)}$$

this is named *E-step* (from "Expectation")

ML and mixtures of gaussians: iterative approach

1. Assign an initial estimate to $\mu_j, \Sigma_j, \pi_j, j = 1, \ldots, K$
2. Repeat
   (a) Compute
   $$\gamma_j(x_i) = \frac{1}{\gamma_i}\pi_j \mathcal{N}(x_i|\mu_j, \Sigma_j) \qquad \text{con} \qquad \gamma_i = \sum_{k=1}^{K}\pi_k \mathcal{N}(x_i|\mu_j, \Sigma_j)$$
   (b) Compute
   $$\pi_j = \frac{n_j}{n} \qquad \text{con} \qquad n_j = \sum_{i=1}^{n}\gamma_j(x_i)$$
   (c) Compute
   $$\mu_j = \frac{1}{n_j}\sum_{i=1}^{n}\gamma_j(x_i)x_i$$
   (d) Compute
   $$\Sigma_j = \frac{1}{n_j}\sum_{i=1}^{n}\gamma_j(x_i)(x_i - \mu_j)(x_i - \mu_j)^T$$
3. until some convergence property is verified

The convergence test may refer to the the increase of log-likelihood in the last iteration

Expectation maximization algorithm

This algorithm is indeed the application of a general schema named *Expectation-Maximization*