



Introduction to Statistical Machine Learning

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(Many figures from C. M. Bishop, "Pattern Recognition and Machine Learning")



Part VII

Mixture Models and EM 1

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



- Sum rule $p(A, B) = \sum_C p(A, B, C)$
- Product rule $p(A, B) = p(A|B)p(B)$
- Why do we optimize the log likelihood?

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



- Estimate best predictor = training = learning

Given data $(x_1, y_1), \dots, (x_n, y_n)$, find a predictor $f_{\mathbf{w}}(\cdot)$.

- 1 Identify the type of input x and output y data
- 2 Propose a (linear) mathematical model for $f_{\mathbf{w}}$
- 3 Design an objective function or likelihood
- 4 Calculate the optimal parameter (\mathbf{w})
- 5 Model uncertainty using the Bayesian approach
- 6 Implement and compute (the algorithm in python)
- 7 Interpret and diagnose results

We will study unsupervised learning this week

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



- Complex **marginal distributions** over observed variables can be expressed via more tractable **joint distributions** over the expanded space of observed and latent variables.
- Mixture Models can also be used to cluster data.
- General technique for finding maximum likelihood estimators in latent variable models:
expectation-maximisation (EM) algorithm.

Review

Motivation

K-means Clustering

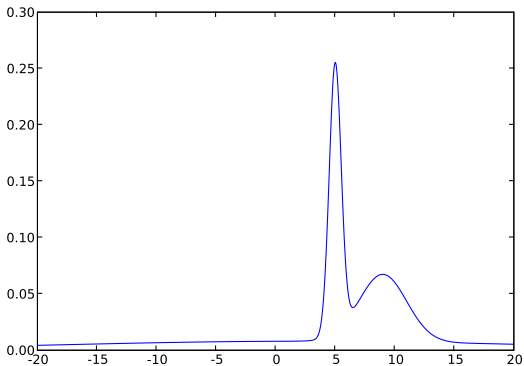
K-means Applications

Mixture of Gaussians

Example - Wallaby Distribution



- Introduced very recently to show ...



Review

Motivation

K-means Clustering

K-means Applications

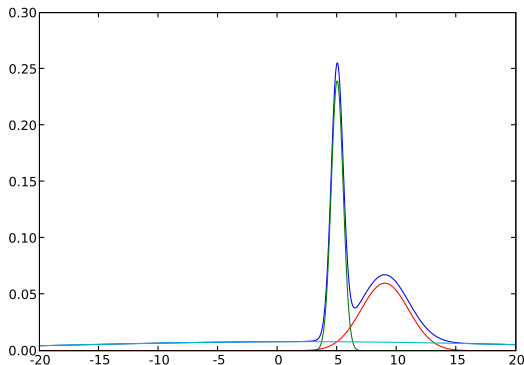
Mixture of Gaussians

Example - 'Wallaby' Distribution



- ... that already a mixture of three Gaussian can be fun.

$$p(x) = \frac{3}{10} \mathcal{N}(x | 5, 0.5) + \frac{3}{10} \mathcal{N}(x | 9, 2) + \frac{4}{10} \mathcal{N}(x | 2, 20)$$



Review

Motivation

K-means Clustering

K-means Applications

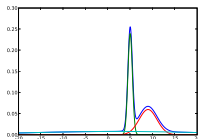
Mixture of Gaussians

Example - 'Wallaby' Distribution

- Use μ, σ as latent variables and define a distribution

$$p(\mu, \sigma) = \begin{cases} \frac{3}{10} & \text{if } (\mu, \sigma) = (5, 0.5) \\ \frac{3}{10} & \text{if } (\mu, \sigma) = (9, 2) \\ \frac{4}{10} & \text{if } (\mu, \sigma) = (2, 20) \\ 0 & \text{otherwise.} \end{cases}$$

$$\begin{aligned} p(x) &= \int_{-\infty}^{\infty} \int_0^{\infty} p(x, \mu, \sigma) \, d\mu \, d\sigma \\ &= \int_{-\infty}^{\infty} \int_0^{\infty} p(x | \mu, \sigma) p(\mu, \sigma) \, d\mu \, d\sigma \\ &= \frac{3}{10} \mathcal{N}(x | 5, 0.5) + \frac{3}{10} \mathcal{N}(x | 9, 2) + \frac{4}{10} \mathcal{N}(x | 2, 20) \end{aligned}$$



Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



- Given a set of data $\{\mathbf{x}_1, \dots, \mathbf{x}_N\}$ where $\mathbf{x}_n \in \mathbb{R}^D$, $n = 1, \dots, N$.
- Goal: Partition the data into K clusters.

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



- Given a set of data $\{\mathbf{x}_1, \dots, \mathbf{x}_N\}$ where $\mathbf{x}_n \in \mathbb{R}^D$, $n = 1, \dots, N$.
- Goal: Partition the data into K clusters.
- Each cluster contains points close to each other.
- Introduce a prototype $\mu_k \in \mathbb{R}^D$ for each cluster.
- Goal: Find
 - 1 a set prototypes μ_k , $k = 1, \dots, K$, each representing a different cluster.
 - 2 an assignment of each data point to exactly one cluster.

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians

K-means Clustering - The Algorithm



Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians

- Start with arbitrary chosen prototypes $\mu_k, k = 1, \dots, K$.
 - 1 Assign each data point to the closest prototype.
 - 2 Calculate new prototypes as the mean of all data points assigned to each of them.
- In the following, we will formalise this introducing a notation which will be useful later.



- Binary indicator variables

$$r_{nk} = \begin{cases} 1, & \text{if data point } \mathbf{x}_n \text{ belongs to cluster } k \\ 0, & \text{otherwise} \end{cases}$$

using the 1-of- K coding scheme.

- Define a **distortion measure**

$$J = \sum_{n=1}^N \sum_{k=1}^K r_{nk} \|\mathbf{x}_n - \boldsymbol{\mu}_k\|^2$$

- Find the values for $\{r_{nk}\}$ and $\{\boldsymbol{\mu}_k\}$ so as to minimise J .

K-means Clustering - Notation

- Find the values for $\{r_{nk}\}$ and $\{\mu_k\}$ so as to minimise J .
- But $\{r_{nk}\}$ depends on $\{\mu_k\}$, and $\{\mu_k\}$ depends on $\{r_{nk}\}$.



K-means Clustering - Notation



- Find the values for $\{r_{nk}\}$ and $\{\mu_k\}$ so as to minimise J .
- But $\{r_{nk}\}$ depends on $\{\mu_k\}$, and $\{\mu_k\}$ depends on $\{r_{nk}\}$.
- Iterate until no further change

- ① Minimise J w.r.t. r_{nk} while keeping $\{\mu_k\}$ fixed,

$$r_{nk} = \begin{cases} 1, & \text{if } k = \arg \min_j \|\mathbf{x}_n - \mu_j\|^2 \\ 0, & \text{otherwise.} \end{cases} \quad \forall n = 1, \dots, N$$

Expectation step

- ② Minimise J w.r.t. $\{\mu_k\}$ while keeping r_{nk} fixed,

$$0 = 2 \sum_{n=1}^N r_{nk} (\mathbf{x}_n - \mu_k)$$
$$\mu_k = \frac{\sum_{n=1}^N r_{nk} \mathbf{x}_n}{\sum_{n=1}^N r_{nk}}$$

Maximisation step

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians

K-means Clustering - Example



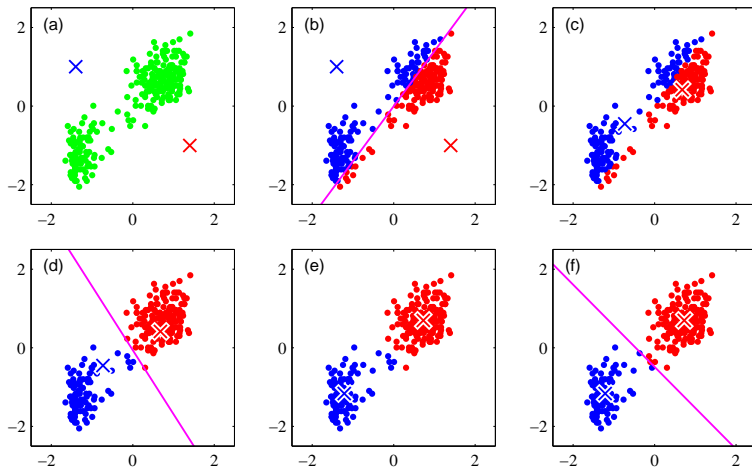
Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



K-means Clustering - Example



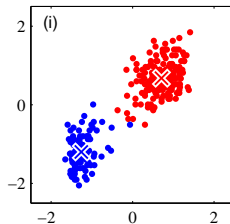
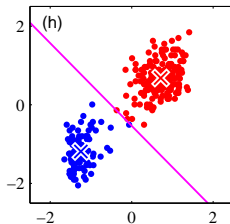
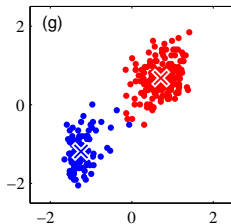
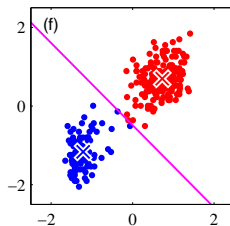
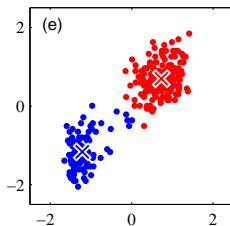
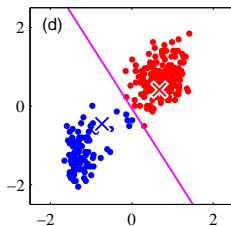
Review

Motivation

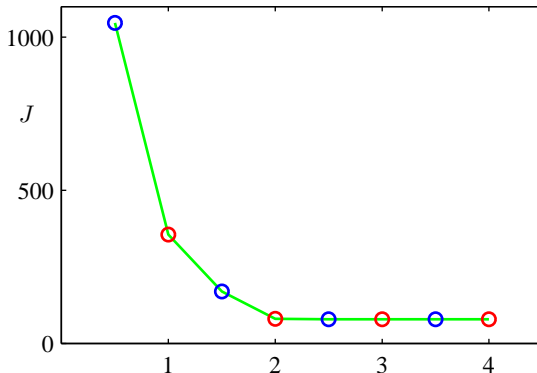
K-means Clustering

K-means Applications

Mixture of Gaussians



K-means Clustering - Cost Function



Cost function J after each E step (blue points)
and M step (red points).

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians

- Initial condition crucial for convergence.
- What happens, if at least one cluster centre is too far from all data points?
- Complex step: Finding the nearest neighbour. (Use triangle inequality; build K-D trees, ...)
- Generalise to non-Euclidean dissimilarity measures $\mathcal{V}(\mathbf{x}_n, \boldsymbol{\mu}_k)$ (called *K-medoids* algorithm),

$$\tilde{J} = \sum_{n=1}^N \sum_{k=1}^K r_{nk} \mathcal{V}(\mathbf{x}_n, \boldsymbol{\mu}_k).$$

- Online stochastic algorithm
 - 1 Draw data point \mathbf{x}_n and locate nearest prototype $\boldsymbol{\mu}_k$.
 - 2 Update only $\boldsymbol{\mu}_k$ using decreasing learning rate η_n

$$\boldsymbol{\mu}_k^{\text{new}} = \boldsymbol{\mu}_k^{\text{old}} + \eta_n (\mathbf{x}_n - \boldsymbol{\mu}_k^{\text{old}}).$$

K-means Clustering - Image Segmentation



- Segment an image into regions of reasonable homogeneous appearance.
- Each pixel is a point in \mathbb{R}^3 (red, blue, green). (Note that the pixel intensities are bounded in the range $[0, 1]$ and therefore this space is strictly speaking not Euclidean).
- Run K -means on all points of the image until convergence. Replace all pixels with the corresponding mean μ_k .
- Results in an image with a palette only K different colours.
- There are much better approaches to image segmentation (but it is an active research topic), this here serves only to illustrate K -means.

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians

Illustrating K-means Clustering - Segmentation



Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians

$K = 2$



$K = 10$



$K = 3$



Original image



Illustrating K-means Clustering - Segmentation

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Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



K-means Clustering - Compression



- **Lossy** data compression: accept some errors in the reconstruction as trade-off for higher compression.
- Apply K -means to the data.
- Store the **code-book vectors** μ_k .
- Store the data in the form of references (labels) to the code-book. Each data point has a label in the range $[1, \dots, K]$.
- New data points are also compressed by finding the closest code-book vector and then storing only the label.
- This technique is also called **vector quantisation**.

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians

Illustrating K-means Clustering - Compression



Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians

$K = 2$



4.2%

$K = 3$



8.3%

$K = 10$



16.7%

Original image



100 %



- We have already seen a mixture of two Gaussians for linear classification
- However in the clustering scenario, we do not observe the class membership
- Strategy (this is vague)
 - We have a difficult distribution $p(\mathbf{x})$
 - We introduce a new variable \mathbf{z} to get $p(\mathbf{x}, \mathbf{z})$
 - Model $p(\mathbf{x}, \mathbf{z}) = p(\mathbf{x}|\mathbf{z})p(\mathbf{z})$ with easy distributions

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



- A **Gaussian mixture distribution** is a linear superposition of Gaussians of the form

$$p(\mathbf{x}) = \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k).$$

- As $\int p(\mathbf{x}) \, d\mathbf{x} = 1$, it follows $\sum_{k=1}^K \pi_k = 1$.
- Let us write this with the help of a **latent** variable \mathbf{z} .

Definition (Latent variables)

Latent variables (as opposed to observable variables), are variables that are not directly observed but are rather inferred (through a mathematical model) from other variables that are observed and directly measured. They are also sometimes called hidden variables, model parameters, or hypothetical variables.

Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



- Let $\mathbf{z} \in \{0, 1\}^K$ and $\sum_{k=1}^K z_k = 1$. In words, \mathbf{z} is a K -dimensional vector in 1-of- K representation.
- There are exactly K different possible vectors \mathbf{z} depending on which of the K entries is 1.
- Define the joint distribution $p(\mathbf{x}, \mathbf{z})$ in terms of a marginal distribution $p(\mathbf{z})$ and a conditional distribution $p(\mathbf{x} | \mathbf{z})$ as

$$p(\mathbf{x}, \mathbf{z}) = p(\mathbf{z}) p(\mathbf{x} | \mathbf{z})$$



Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



- Set the marginal distribution to

$$p(z_k = 1) = \pi_k$$

where $0 \leq \pi_k \leq 1$ together with $\sum_{k=1}^K \pi_k = 1$.

- Because \mathbf{z} uses 1-of- K coding, we can also write

$$p(\mathbf{z}) = \prod_{k=1}^K \pi_k^{z_k}.$$

- Set the conditional distribution of \mathbf{x} given a particular \mathbf{z} to

$$p(\mathbf{x} | z_k = 1) = \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k),$$

or

$$p(\mathbf{x} | \mathbf{z}) = \prod_{k=1}^K \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)^{z_k},$$



- The marginal distribution over \mathbf{x} is now found by summing the joint distribution over all possible states of \mathbf{z}

$$\begin{aligned} p(\mathbf{x}) &= \sum_{\mathbf{z}} p(\mathbf{z}) p(\mathbf{x} | \mathbf{z}) = \sum_{\mathbf{z}} \prod_{k=1}^K \pi_k^{z_k} \prod_{k=1}^K \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)^{z_k} \\ &= \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \end{aligned}$$

- The marginal distribution of \mathbf{x} is a Gaussian mixture.
- For several observations $\mathbf{x}_1, \dots, \mathbf{x}_N$ we need one latent variable \mathbf{z}_n per observation.
- What have we gained? Can now work with the joint distribution $p(\mathbf{x}, \mathbf{z})$. Will lead to significant simplification later, especially for EM algorithm.



Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians

- Conditional probability of \mathbf{z} given \mathbf{x} by Bayes' theorem

$$\begin{aligned}\gamma(z_k) = p(z_k = 1 | \mathbf{x}) &= \frac{p(z_k = 1) p(\mathbf{x} | z_k = 1)}{\sum_{j=1}^K p(z_j = 1) p(\mathbf{x} | z_j = 1)} \\ &= \frac{\pi_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)}\end{aligned}$$

- $\gamma(z_k)$ is the **responsibility** of component k to 'explain' the observation \mathbf{x} .

Mixture of Gaussians - Ancestral Sampling



- Goal: Generate random samples distributed according to the mixture model.
 - 1 Generate a sample $\hat{\mathbf{z}}$ from the distribution $p(\mathbf{z})$.
 - 2 Generate a value $\hat{\mathbf{x}}$ from the conditional distribution $p(\mathbf{x} | \hat{\mathbf{z}})$.
- Example: Mixture of 3 Gaussians, 500 points.

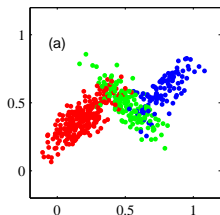
Review

Motivation

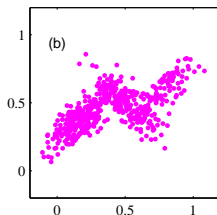
K-means Clustering

K-means Applications

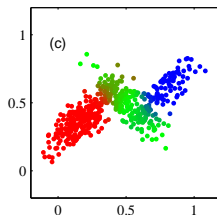
Mixture of Gaussians



Original states of \mathbf{z} .



Marginal $p(\mathbf{x})$.

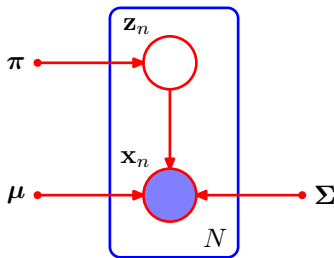


(R, G, B) - colours
mixed according to
 $\gamma(z_{nk})$.

Mixture of Gaussians - Maximum Likelihood



- Given N data points, each of dimension D , we have the data matrix $\mathbf{X} \in \mathbb{R}^{N \times D}$ where each row contains one data point.
- Similarly, we have the matrix of latent variables $\mathbf{Z} \in \mathbb{R}^{N \times K}$ with rows \mathbf{z}_n^T .
- Assume the data are drawn i.i.d., the distribution for the data can be represented by a graphical model.



Review

Motivation

K-means Clustering

K-means Applications

Mixture of Gaussians



- The log of the likelihood function is then

$$\ln p(\mathbf{X} | \boldsymbol{\pi}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \sum_{n=1}^N \ln \left\{ \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right\}$$

- Significant problem: If a mean $\boldsymbol{\mu}_j$ 'sits' directly on a data point \mathbf{x}_n then

$$\mathcal{N}(\mathbf{x}_n | \mathbf{x}_n, \sigma_j^2 \mathbf{I}) = \frac{1}{(2\pi)^{1/2}} \frac{1}{\sigma_j}.$$

- Here we assumed $\boldsymbol{\Sigma}_k = \sigma_k^2 \mathbf{I}$. But problem is general, just think of a main axis transformation for $\boldsymbol{\Sigma}_k$.
- Overfitting (in disguise) occurring again with the maximum likelihood approach.
- Use heuristics to detect this situation and reset the mean of the corresponding component of the mixture.

Mixture of Gaussians - Maximum Likelihood



- A K component mixture has a total of $K!$ equivalent solutions corresponding to the $K!$ ways of assigning K sets of parameters to K solutions.
- Also called **identifiability problem**. Needs to be considered when the parameters discovered by a model are interpreted.
- Maximising the log likelihood of a Gaussian mixture is more complex than for a single Gaussian. Summation over all K components inside of the logarithm make it harder.
- Setting the derivatives of the log likelihood to zero does not longer result in a closed form.
- May use gradient-based optimisation.
- Or EM algorithm. Stay tuned.