

# IAML DL - Study Guide - Week 07

Sambit Paul, Pavlos Andreadis

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## 1 Introduction

Week 7 introduces 2 unsupervised learning methods for clustering of data.

The first method being explored is K-means which aims to cluster data into  $K$  groups by minimizing a criterion known as *inertia*.  $K$  is a parameter that needs to be chosen as a parameter before execution by the user.

Along side that, we will also explore Gaussian mixture models (GMMs) which are a generalisation of K-means to incorporate covariance information [Pedregosa et al. \[2011\]](#). This model uses a combination of Gaussian distributions to model the data.

## 2 K-Means Clustering

- Why is it called **K-Means**?  
In K-Means the term  $K$  refers to the number of clusters that need to be identified; and, *means* refers to the process of averaging of data to find the centroid of each cluster.
- **Monothetic and Polythetic Clustering:** In a monothetic scheme, cluster membership is based on the presence or absence of a single characteristic. Polythetic schemes use more than one characteristic. For example, classifying people solely on the basis of their gender is a monothetic classification, but if both gender and handedness (left or right handed) are used, the classification is polythetic.
- To read about hard and soft clustering, please refer to this [article](#).
- The objective of K-means as defined in [Bishop \[2006\]](#) Section 9.1 is the minimisation of the cost function  $J$  where  $J = \sum_{n=1}^N \sum_{k=1}^K r_{n,k} \|x_n - \mu_k\|^2$  such that,  $r_{n,k}$  denotes if point  $n$  belongs to cluster  $k$  and  $\|x_n - \mu_k\|^2$  is the squared error.
- To understand the K-means algorithm, please refer to [Wu et al. \[2008\]](#) Section 2.1. The basic steps can be elucidated as:

1. Specify number of clusters  $K$ .
  2. Initialize centroids by first shuffling the dataset and then randomly selecting  $K$  data points for the centroids without replacement.
  3. Keep iterating until there is no change to the centroids or maximum iterations has been reached.
- An improvement on the basic K-means algorithm is to introduce a kernel on top of the data to project it into a high-dimensional space [Dhillon et al. \[2004\]](#). Although the boundaries will be linear in the high-dimensional space, on projecting back to the lower dimensions, it becomes non-linear.
  - To read about the limitations of K-means, please refer to [Wu et al. \[2008\]](#) Section 2.2.
  - To get a quick overview of the K-means algorithm, please refer to [Barber \[2012\]](#) Section 20.3.5. [Requires an understanding of Expectation Maximization]

### 3 Gaussian Mixture Models

- This topic requires an intuition about Maximum Likelihood Estimation. To get a quick refresher, please refer to this [article](#).
- What is **Expectation-Maximization**?  
Expectation maximization is an iterative process of improving the probability of a model to predict if an observation belongs to a specific distribution in the presence of latent variables.
  - E-Step  $\Rightarrow$  Estimate the missing variables in the dataset
  - M-Step  $\Rightarrow$  Maximize the parameters of the model in the presence of the data

Maximum Likelihood estimate the same probability in the absence of latent variables.

- [This](#) can be used good starter video to understand the intuition about Expectation-Maximization (EM).
- To get a deeper understanding of the mathematics behind the general EM algorithm, please refer to [Bishop \[2006\]](#) Section 9.4. Another approach to EM, based on mathematical derivations, is provided in Section 2 of this [document](#).
- Basic Representation of Mixture Models is provided in [Figure 1](#)
- An intuitive concept of Gaussian Mixture model is provided in this [article](#)

<b>Data:</b>	$\mathcal{D} = \{\mathbf{x}^{(i)}\}_{i=1}^N$ where $\mathbf{x}^{(i)} \in \mathbb{R}^M$
<b>Generative Story:</b>	$z \sim \text{Categorical}(\phi)$ $\mathbf{x} \sim p_{\theta}(\cdot z)$
<b>Model:</b>	Joint: $p_{\theta, \phi}(\mathbf{x}, z) = p_{\theta}(\mathbf{x} z)p_{\phi}(z)$ Marginal: $p_{\theta, \phi}(\mathbf{x}) = \sum_{z=1}^K p_{\theta}(\mathbf{x} z)p_{\phi}(z)$
<b>(Marginal) Log-likelihood:</b>	$\ell(\theta) = \log \prod_{i=1}^N p_{\theta, \phi}(\mathbf{x}^{(i)})$ $= \sum_{i=1}^N \log \sum_{z=1}^K p_{\theta}(\mathbf{x}^{(i)} z)p_{\phi}(z)$

Figure 1: These are the basic steps that need to be followed to build a Mixture Model

- Section 2 and 3 from this [document](#) provides an elaborate explanation of Gaussian Mixture models and Expectation Maximization.
- A thorough and clear explanation of Gaussian Mixture Models (albeit, slightly lengthy) is also provided in [Bishop \[2006\]](#) Section 9.2.

## 4 Comparison between K-means and GMM

Criterion	K-Means	GMM
<i>Convergence</i>	Faster than GMM	Slower than K-Means
<i>Speed</i>	Computationally less intensive	Computationally intensive
<i>Initialization</i>	Random Initialisation	Use K-means to determine the means of the Gaussian
<i>Output</i>	Single hard assignment to clusters	Probability distribution over the cluster assignment

Table 1: This table provides a comparative analysis of K-Means clustering and Gaussian Mixture Models over 4 criteria

## References

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