IAML DL - Study Guide - Week 7

Sambit Paul, Pavlos Andreadis, Nigel Goddard

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1 Introduction

Week 7 introduces 2 unsupervised learning methods for clustering of data. The first method being explored in 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

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

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
Convergence	Faster than GMM	Slower than K-Means
Speed	Computationally less	Computationally in-
	intensive	tensive
Initialization	Random Initialisation	Use K-means to deter-
		mine the means of the
		Gaussian
Output	Single hard assignment	Probability distribu-
	to clusters	tion 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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