

ST451 - Lent term

Bayesian Machine Learning

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Clusters/Mixture Models and EM algorithm

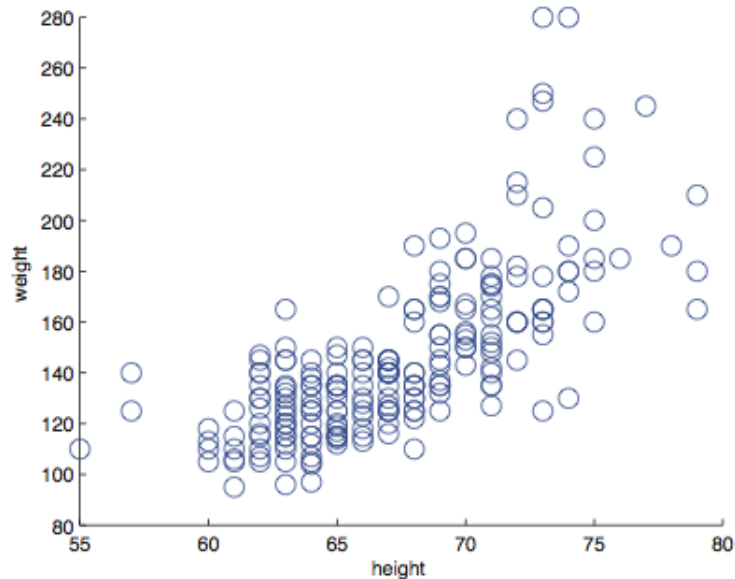
Outline

- 1 Introduction
- 2 Mixture models
- 3 EM algorithm
- 4 Fully Bayesian mixture models

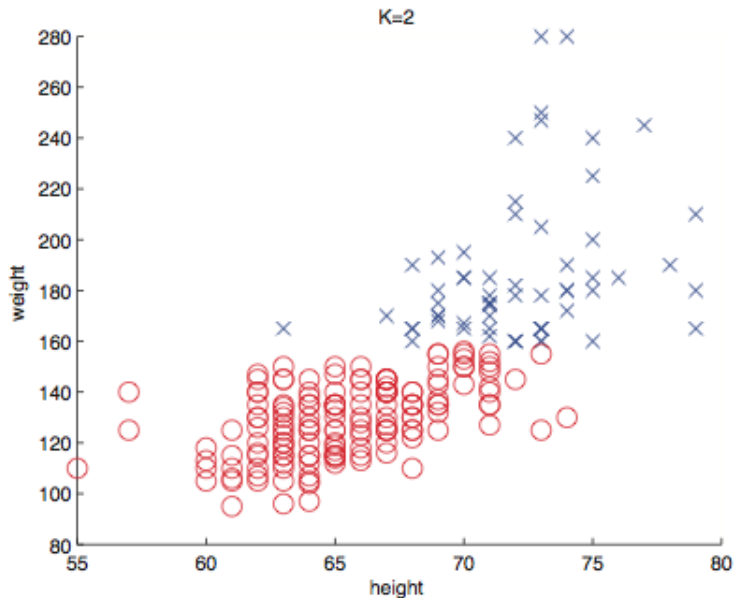
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Motivating Example 1: Heights and weights



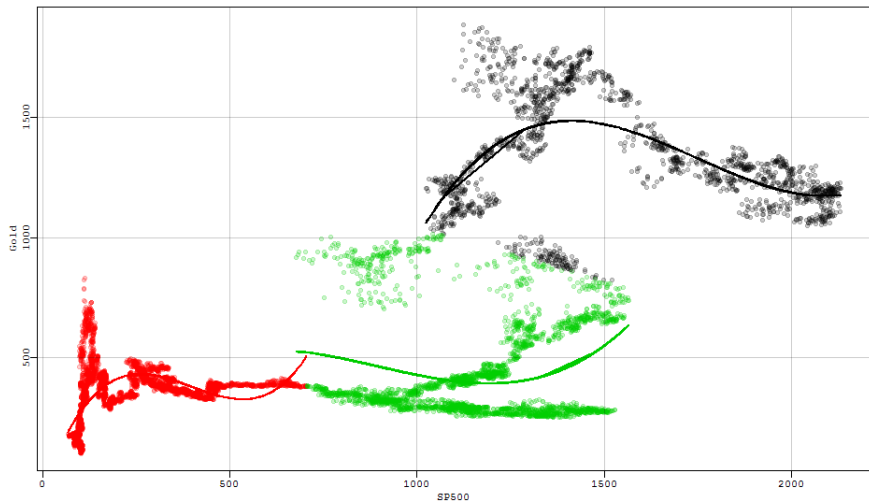
Example 1: Heights and weights



Example 2: Job Quality Definition/Measurement



Example 3: Financial Modelling



Example 4: Image Compression

$K = 2$



$K = 3$



$K = 10$



Original image

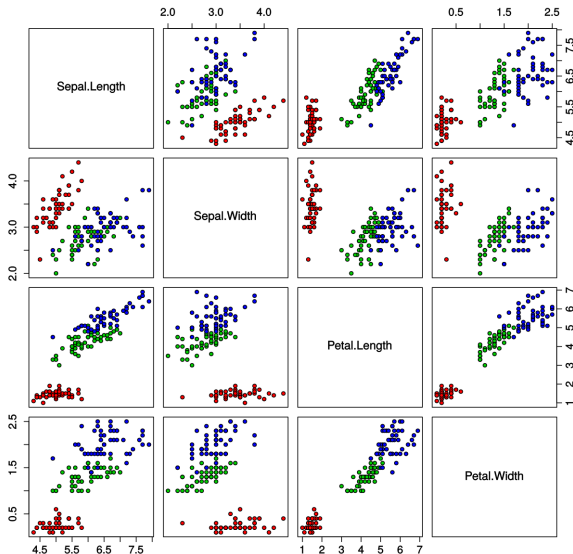


Example 5: Distinguishing Iris flower species



Example 5: Distinguishing Iris flower species

Iris Data (red=setosa,green=versicolor,blue=virginica)



Example 6: Bayesian non-parametric density models

Recall the **VIX** index and the model we used to capture some of its **stylised facts**

$$Y_t = Y_{t-1} + \kappa(\mu - Y_{t-1})\delta + \sigma\epsilon_t,$$

where Y_t is VIX at time t , δ is the time interval, and ϵ_t are **independent** error terms.

The distribution of each ϵ_t may assumed to be a **mixture** of Normal distributions.

Such model is very **flexible**; in this case corresponds to a **jump diffusion** model.

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Data augmentation

Often we want to draw inference on parameters θ based on data x from a likelihood $f(x|\theta)$ that is either **intractable or expensive to compute**.

Introduce an **unobserved latent variable** z to extend the model defining $f(z, x|\theta)$

We can then work **directly** with $f(z, x|\theta)$ (variational Bayes, MCMC) or approximate the integral $f(x|\theta) = \int f(z, x|\theta) dz$ in some way (simulated likelihood, EM).

Many **famous examples**, e.g. Ising model, factor analysis, random effects, hidden Markov models and mixtures.

Cluster/mixture analysis

- The populations consists of K **clusters/groups**, each with distribution $f(x_i|\theta_k)$, $k = 1, \dots, K$.
- Each individual $i = 1, \dots, n$, belongs to **one** of these K clusters.
- Cluster indicator $z_i = 1, \dots, K$ is an **unobserved/latent** categorical variable with Multinoulli distribution $\pi(z|\pi_k)$, where $\sum_k \pi_k = 1$.
- The aim is to **classify** individuals, by and draw **inference** on $\theta = (\pi_k, \theta_k)_{k=1}^K$.

Likelihood and augmented likelihood

Define also the z_{ik} indicator that takes the value 1 if the individual i is in cluster k and 0 otherwise. So if $z_i = 2$, $z_{i2} = 1$ and $z_{ik} = 0$ for $k \neq 2$.

The **augmented likelihood** also includes z_i for each x_i .

$$f(z_i, x_i | \theta) = \pi(z_i | \pi_k) f(x_i | z_i, \theta_k) = \prod_{k=1}^K \pi_k^{z_{ik}} f(x_i | \theta_k)^{z_{ik}}.$$

Note that $f(z_i = k, x_i | \theta) = \pi_k f(x_i | \theta_k)$. **Summing out** z_i gives

$$f(x_i | \theta) = \sum_{k=1}^K f(z_i = k, x_i | \theta) = \sum_{k=1}^K \pi_k f(x_i | \theta_k)$$

Overall we have $f(x | \theta) = \prod_{i=1}^n f(x_i | \theta)$ and $f(z, x | \theta) = \prod_{i=1}^n f(x_i, z_i | \theta)$

Example: Gaussian Mixture Models

In **Gaussian mixture models**, we have $x_i|z = k \sim N(\mu_k, \Sigma_k)$

Hence

$$f(x_i|\theta) = \sum_{k=1}^K \pi_k N(x_i|\mu_k, \Sigma_k)$$

so the **parameters** to be estimated are $\theta = (\pi_k, \mu_k, \Sigma_k)_{k=1}^K$.

Due to the **large number** of parameters, especially for large K , restrictions are often placed on Σ_k , e.g. diagonal or tied.

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Main idea

Complete Data: If we knew the cluster each person is, z_i , then MLE is straightforward: split the data into clusters do MLE in each cluster separately.

But we don't, so we need a modified approach. The algorithm used most frequently is the **EM**.

A rough sketch is the one below

- 1 Start with a θ .
- 2 **E step:** Use Bayes theorem to find the **responsibilities** $\gamma_{ik} = \pi(z_i = k|x, \theta)$ to get the **expected** log likelihood.
- 3 **M step:** **Maximise** the expected log-likelihood and update θ .
- 4 Continue until convergence.

log-likelihood and augmented log-likelihood

First write down the **augmented log-likelihood**. Remember that

$$f(z_i, x_i | \theta) = \prod_{k=1}^K \pi_k^{z_{ik}} f(x_i | \theta_k)^{z_{ik}},$$

so considering all individuals and taking log gives

$$\log f(z, x | \theta) = \sum_{i=1}^n \sum_{k=1}^K z_{ik} (\log \pi_k + \log f(x_i | \theta_k))$$

By contrast the **log-likelihood** is

$$\log f(x | \theta) = \log \left[\sum_{k=1}^K \pi_k f(x_i | \theta_k) \right]$$

Notes

- 1 Can view the augmentation as way to bring log within the sum.
- 2 Easy to maximise the augmented log-likelihood given the z_{ik} 's.

E step

In the EM algorithm we update θ^{old} to θ^{new} . In the **E step** we define the **expected log likelihood**

$$Q(\theta, \theta^{old}) = \mathbb{E}_{\pi(z|x, \theta^{old})} \left[\sum_{i=1}^n \sum_{k=1}^K z_{ik} (\log \pi_k + \log f(x_i | \theta_k)) \right]$$

Note that

$$\mathbb{E}_{\pi(z|x, \theta^{old})} [z_{ik}] = \frac{\pi_k^{old} f(x_i | \theta_k^{old})}{\sum_{j=1}^K \pi_j^{old} f(x_i | \theta_j^{old})} = \gamma(z_{ik}),$$

where the $\gamma(z_{ik}) = \pi(z_{ik} = 1 | x, \theta^{old})$ are known as the **responsibilities**.

Hence we can write

$$Q(\theta, \theta^{old}) = \sum_{i=1}^n \sum_{k=1}^K \gamma(z_{ik}) (\log \pi_k + \log f(x_i | \theta_k))$$

M step

The **M step**: consists of simply maximising $Q(\theta, \theta^{old})$ wrt to θ . Note that the $\gamma(z_{ik})$ are known numbers based on x and θ^{old} so it is usually an easy task.

To maximising $Q(\theta, \theta^{old})$ wrt to π_k 's we can use **Lagrange multipliers** to satisfy the restriction that they sum to one. So we set

$$L = Q(\theta, \theta^{old}) + \lambda \left(\sum_k \pi_k - 1 \right),$$
$$\frac{\partial L}{\partial \pi_k} = 0 \quad \Leftrightarrow \quad \pi_k = \frac{\sum_i \gamma(z_{ik})}{-\lambda}$$
$$\frac{\partial L}{\partial \lambda} = 0 \quad \Leftrightarrow \quad \sum_k \pi_k = 1 \Leftrightarrow \lambda = -n$$

so we get that $Q(\theta, \theta^{old})$ is maximised at

$$\pi_k^{new} = \frac{\sum_i \gamma(z_{ik})}{n} = \frac{n_k}{n}$$

Example: Gaussian Mixture models

The **remaining** parameters depend on which type of $f(x_i|\theta_k)$ we have.

For Gaussian mixture models standard **MLE** methods provide

$$\begin{aligned}\mu_k^{new} &= \frac{\sum_i \gamma(z_{ik}) x_i}{\sum_i \gamma(z_{ik})} = \frac{\sum_i \gamma(z_{ik}) x_i}{n_k} \\ \Sigma_k^{new} &= n \frac{1}{n_k} \sum_i \gamma(z_{ik}) (x_i - \mu_k^{new})(x_i - \mu_k^{new})^T\end{aligned}$$

Hence the EM algorithm **initiates** θ and iteratively **updates** from θ^{old} to θ^{new} until the log likelihood or the parameters converge.

Similar results exist for **other** distributions such as Bernoulli, Exponential etc.

Connection with k-means

- Mixture models classify individuals to clusters based on the **responsibilities** $\gamma(\zeta_{ik})$'s, i.e. the posterior probabilities of z , rather than with certainty, aka **soft allocation**.
- This is reflected on the estimate of θ_k that are **weighted averages** based on how likely an individual is in cluster k .
- In Gaussian mixture models if we set $\Sigma_k = \sigma^2 I_d$ and let $\sigma^2 \rightarrow 0$ we get the same solution as with the k-means approach for μ_k . Note that in this case we have **hard allocation**.
- If we have general Σ'_k s the approach coincide with the **elliptical** k-means.

Selecting the number of clusters

- In both mixture models and k-means it is **not easy** to select the number of classes.
- The default criterion in the mixture models is the **BIC**.
- Nevertheless the approach is very sensitive to starting values as the objective is multimodal and is very likely to get trapped in **local maxima**.
- It is recommended to initialise parameters based on intuition, try out multiple starting points or initialise with the results of another method.

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Fully Bayesian approach

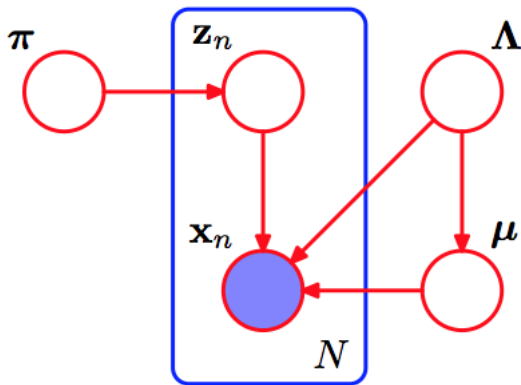
The approach so far was Bayesian with respect to z but **not** θ . For a fully Bayesian approach priors on θ should be specified.

In Gaussian mixture models example the **conjugate priors** can be used

$$\begin{aligned}\mu_k | \Lambda_k &\sim N\left(m_0, (\beta_0 \Lambda)^{-1}\right) \\ \Lambda &\sim \text{Wishart}(W_0, \nu_0) \\ \pi &\sim \text{Dirichlet}(\alpha_0)\end{aligned}$$

The posterior is not available in closed form. We can therefore consider a **variational approximation**.

Graph of Bayesian Gaussian Mixtures Model



Variational Inference for Gaussian Mixtures

We can apply **mean field** approximation of the form

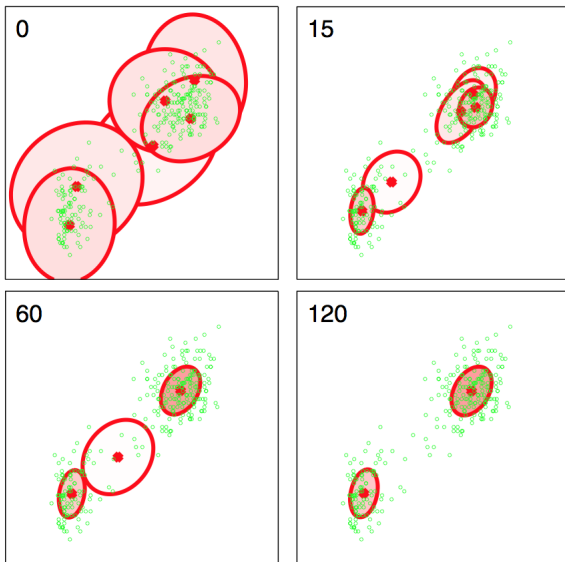
$$q(z, \pi, \mu, \Lambda) = q(z)q(\pi, \mu, \Lambda)$$

The outcome is analogous to the EM case.

- 1 $q(z)$ requires some tedious calculations that provide the responsibilities r_{ik} 's
- 2 Given r_{ik} 's $q(\pi, \mu, \Lambda)$ **can be recognised** as a product of a Dirichlet, Normal and Wishart distributions, the moments of which can be used to update r_{ik} 's. See Bishop 478-79 for details.
- 3 The above are repeated until convergence.

The prior parameter α_0 , in the Dirichlet(α_0) prior on π , is **critical** in selecting the number of clusters. Large values favour equal π_k 's but under values close to 0 only the important clusters will get individuals.

Variational Inference - no overfit



Latent Dirichlet allocation

Consider M documents each with N_i words. The model is defined as:

- Each of word w_{ij} , $i = 1, \dots, M$ and $j = 1, \dots, N_i$ may belong to different topics $k = 1, \dots, K$, that indicate the word distribution ϕ_k :

$$w_{ij}|z_{ij} \sim \text{Multinoulli}(\phi_k)$$

- Let z_{ij} denote the topic indicator (latent) of w_{ij} , with distribution θ_i varying across documents:

$$z_{ij} \sim \text{Multinoulli}(\theta_i)$$

- The word and topic distributions (θ_i, ϕ_k) are given Dirichlet distribution priors with α, β respectively, both < 1 for sparsity:

$$\theta_i \sim \text{Dirichlet}(\alpha) \quad \phi_k \sim \text{Dirichlet}(\beta)$$

Today's lecture - Reading

Bishop: 9.1 to 9.4, 10.2.1 10.2.2

Murphy: 11.1 11.2 11.4.1 11.4.2 21.6