Expectation Maximization

Maschinelles Lernen 1 -Grundverfahren WS20/21

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Learning Outcomes

What will we learn today?

- Understand latent variable models and why they are hard to train
- Understand mixture models and how to train it using EM
- Analysis of the EM algorithm and why it converges

Agenda for today

Mixture Models

- Gaussian Mixture Models (GMM)
- Expectation Maximization

Latent Variable Models and Generalized EM

- EM decomposition
- E- and M-step
- Convergence analysis
- EM for dimensionality reduction

Mixture Models

Parametric models

- Gaussian, Neural Networks, ...
- ✓ Good analytic properties
- ✓ Simple
- ✓ Small memory requirements
- ✓ Fast
- Limited representation power (most parametric distributions have only one mode)

Non-Parametric models

- Kernel-density estimation, k-NN
- ✓ General (can represent any distribution)
- × Curse of dimensionality
- × Large memory requirements
- × Slow

- Mixture models combine the advantages of both worlds
- Key idea: Create a complex distribution by combining simple ones (e.g. Gaussians)

Mixture model





• In the limit with many / infinite components, this can approximate any smooth density

Example: Mixture of Gaussians (MoG) Individual Gaussians • p(x)xSum of Gaussians p(x)x

Gaussian Mixture Models (GMMs)

• Mixture coefficient:

 $p(k) = \pi_k$, with $0 \le \pi_k \le 1$, $\sum_{i} \pi_k = 1$

• Mixture component:

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

• Mixture distribution:

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

- Always integrates to 1
- Parameters of the mixture

$$\boldsymbol{\theta} = \{\pi_1, \boldsymbol{\mu}_1, \boldsymbol{\Sigma}_1, \dots, \pi_K, \boldsymbol{\mu}_K, \boldsymbol{\Sigma}_K\}$$

Example: Mixture of Gaussians (MoG)

Individual Gaussians



Maximum Likelihood of a mixture

(Marginal-)Log-Likelihood with N iid. points

$$\mathcal{L} = \log L(\boldsymbol{\theta}) = \sum_{i=1}^{N} \log \underbrace{p_{\boldsymbol{\theta}}(\boldsymbol{x}_i)}_{\text{marginal}} = \sum_{i=1}^{N} \log \underbrace{\left(\sum_{k=1}^{K} \pi_k \mathcal{N}(\boldsymbol{x}_i | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)\right)}_{\text{non-exponential family}}$$

• Q: Can we do gradient descent?

$$egin{aligned} rac{\partial \mathcal{L}}{\partial oldsymbol{\mu}_j} &= \sum_{i=1}^N rac{\pi_j \mathcal{N}(oldsymbol{x}_i | oldsymbol{\mu}_j, oldsymbol{\Sigma}_j)}{\sum_{k=1}^K \pi_k \mathcal{N}(oldsymbol{x}_i | oldsymbol{\mu}_k, oldsymbol{\Sigma}_k)} oldsymbol{\Sigma}_j^{-1}(oldsymbol{x}_i - oldsymbol{\mu}_j) \ &= \sum_{i=1}^N rac{p(j) p(oldsymbol{x}_i | j)}{p(oldsymbol{x}_i)} oldsymbol{\Sigma}_j^{-1}(oldsymbol{x}_i - oldsymbol{\mu}_j) \ &= \sum_{i=1}^N oldsymbol{\Sigma}_j^{-1}(oldsymbol{x}_i - oldsymbol{\mu}_j) p(j | oldsymbol{x}_i) \end{aligned}$$

- × Gradient depends on all other components (cyclic dependency)
- × No closed form solution
- × Typically very slow convergence
- A: Yes, but the sum (marginalization) does not go well with the log

Estimating Gaussian Mixture Models

So why is optimizing
$$\mathcal{L} = \sum_{i=1}^{N} \log p_{\theta}(\boldsymbol{x}_i)$$
 so hard?

- Because we do not know which mixture component k created which data-point
- If we would have data from the joint distribution $p(\boldsymbol{x}_i, k_i | \boldsymbol{\theta})$, then it would be easy ...

In this case, we can simply perform a maximum likelihood estimate:

• Coefficients:
$$\pi_k = \frac{\sum_i q_{ik}}{N}$$

• Means: $\boldsymbol{\mu}_k = \frac{\sum_i q_{ik} \boldsymbol{x}_i}{\sum_i q_{ik}}$
• Covariances: $\boldsymbol{\Sigma}_k = \frac{\sum_i q_{ik} (\boldsymbol{x}_i - \boldsymbol{\mu}_k) (\boldsymbol{x}_i - \boldsymbol{\mu}_k)^T}{\sum_i q_{ik}}$

where $q_{ik} = \mathbb{I}(k, k_i)$ is 1 if the ith sample belongs to the kth component and 0 otherwise

Expectation-Maximization: Chicken and Egg...

Yet, we do not know which component belongs to which sample

- Can we estimate that? Given a current mixture model, yes!
- Expectation Step:
 - Compute cluster probabilities aka responsibilities for each sample (Bayes rule)

$$q_{ik} = p(k_i = k | \boldsymbol{x}_i) = \frac{p(\boldsymbol{x}_i | k) p(k)}{\sum_j p(\boldsymbol{x}_i | j) p(j)} = \frac{\pi_k \mathcal{N}(\boldsymbol{x}_i | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(\boldsymbol{x}_i | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)}$$

- Responsibilities q_{ik} are now continues between 0 and 1
- But we need to know the Gaussian components

Expectation-Maximization: Chicken and Egg...

Yet, we do not know which component belongs to which sample

- Can we estimate that? Given a current mixture model, yes!
- Maximization Step:
 - Compute (weighted) maximum likelihood estimate

$$\pi_k = \frac{\sum_i q_{ik}}{N} \qquad \qquad \boldsymbol{\mu}_k = \frac{\sum_i q_{ik} \boldsymbol{x}_i}{\sum_i q_{ik}} \qquad \qquad \boldsymbol{\Sigma}_k = \frac{\sum_i q_{ik} (\boldsymbol{x}_i - \boldsymbol{\mu}_k) (\boldsymbol{x}_i - \boldsymbol{\mu}_k)^T}{\sum_i q_{ik}}$$

- But we need to know the responsibilities

Algorithm: EM for GMMs

- **Initialize:** Mixture Components + Mixture coefficients
 - E.g. Use k-means for the component means and some initial covariance
- Repeat until convergence:
 - Expectation-step: Compute responsibilities

$$q_{ik} = \frac{\pi_k \mathcal{N}(\boldsymbol{x}_i | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(\boldsymbol{x}_i | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)}$$

- Maximization-step: Update coefficients, components means and component variance

$$\pi_k = \frac{\sum_i q_{ik}}{N} \qquad \boldsymbol{\mu}_k = \frac{\sum_i q_{ik} \boldsymbol{x}_i}{\sum_i q_{ik}} \qquad \boldsymbol{\Sigma}_k = \frac{\sum_i q_{ik} (\boldsymbol{x}_i - \boldsymbol{\mu}_k) (\boldsymbol{x}_i - \boldsymbol{\mu}_k)^T}{\sum_i q_{ik}}$$

Illustration

- Each component represents a cluster in the data set
- EM is very sensitive to the initialization



EM versus k-means

K-means can be seen as special case of EM with:

- Co-variances are always set to 0 (in the limit)
- E-Step / Assignment Step:
 - responsibilities q_{ik} of nearest cluster k are set to 1, all other values are 0
- M-Step / Adjustment Step:
 - Update for the mean is the same
 - Co-Variances are ignored (set to close to 0)
- EM is harder to learn than k-means but it also gives you variances and densities
- Often k-means is used to initialize the means for EM

K-means is known to converge, does also EM always converge?

• Mixture coefficient:

 $p(k) = \pi_k$, with $0 \le \pi_k \le 1$, $\sum_{k=1}^{\infty} \pi_k = 1$

• Mixture component:

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

• Mixture distribution:

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

- Always integrates to 1
- Parameters of the mixture

$$\boldsymbol{\theta} = \{\pi_1, \boldsymbol{\mu}_1, \boldsymbol{\Sigma}_1, \dots, \pi_K, \boldsymbol{\mu}_K, \boldsymbol{\Sigma}_K\}$$

E-Step

Compute "responsibilities"

$$q_{ik} = \frac{\pi_k \mathcal{N}(\boldsymbol{x}_i | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(\boldsymbol{x}_i | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)} = p(z = k | \boldsymbol{x}_i)$$

 How much component k contributes to generation of x_i according to current mixture model

• Mixture coefficient:

 $p(k) = \pi_k$, with $0 \le \pi_k \le 1$, $\sum_k \pi_k = 1$

• Mixture component:

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

• Mixture distribution:

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

- Always integrates to 1
- Parameters of the mixture

$$\boldsymbol{\theta} = \{\pi_1, \boldsymbol{\mu}_1, \boldsymbol{\Sigma}_1, \dots, \pi_K, \boldsymbol{\mu}_K, \boldsymbol{\Sigma}_K\}$$

M-Step: $\boldsymbol{\theta} = \operatorname*{arg\,max}_{\boldsymbol{\theta}} \sum_{i} \sum_{k} q_{ik} \log p(k) p(\boldsymbol{x}_i | k)$

- We can separate updates of single components and coefficients
 - just additive objectives in lower bound
- Update coefficients:

$$\pi = rg\max_{\pi} \sum_{i} \sum_{k} q_{ik} \log \pi_k$$

Update components:

$$oldsymbol{\mu}_k, oldsymbol{\Sigma}_k = rgmax_{oldsymbol{\mu}_k} \sum_i q_{ik} \log \mathcal{N}(oldsymbol{x}_i | oldsymbol{\mu}_k, oldsymbol{\Sigma}_k)$$

- Each data-point is weighted by q_{ik}
- Weighted maximum likelihood estimate

Weighted Maximum Likelihood updates:

- Update coefficients: $\pi = \arg \max_{\pi} \sum_{i} \sum_{k} q_{ik} \log \pi_k$ - Result: $\pi_k = \frac{\sum_i q_{ik}}{\sum_k \sum_i q_{ik}} = \frac{\sum_i q_{ik}}{N}$
- Update components: $\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k = \operatorname*{arg\,max}_{\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k} \sum_i q_{ik} \log \mathcal{N}(\boldsymbol{x}_i | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)$

- Mean:
$$\boldsymbol{\mu}_k = rac{\sum_i q_{ik} \boldsymbol{x}_i}{\sum_i q_{ik}}$$

- Covariance:
$$\mathbf{\Sigma}_k = rac{\sum_i q_{ik} (\mathbf{x}_i - \mathbf{\mu}_k) (\mathbf{x}_i - \mathbf{\mu}_k)^T}{\sum_i q_{ik}}$$

Agenda for today

Mixture Models

Gaussian Mixture Models (GMM)

The Expectation Maximization (EM) Algorithm

- EM decomposition
- E- and M-step
- Convergence analysis
- EM for GMMs

Latent Variable Models and a generalized view on EM

Mixture Models are Latent Variable Models

Mixture models are an instance of latent variable models

- Examples: mixture models, missing data, latent factors,
- Observed variables: x, Latent variables: z (e.g., index of mixture component) •
- $p_{\boldsymbol{\theta}}(\boldsymbol{x}, \boldsymbol{z})$ Parametric model: ٠
- Marginal distribution: .

$$p_{\theta}(\boldsymbol{x}) = \sum_{z} p(\boldsymbol{x}, z), \quad p_{\theta}(\boldsymbol{x}) = \int_{\boldsymbol{z}} p_{\theta}(\boldsymbol{x}, \boldsymbol{z}) d\boldsymbol{z}$$

discrete latent variable continuous latent variable

 $p_{\boldsymbol{\theta}}(\boldsymbol{x}, \boldsymbol{z}) d\boldsymbol{z}$

(Marginal) Log-Likelihood:

$$L(\boldsymbol{\theta}) = \sum_{i=1}^{N} \log p_{\boldsymbol{\theta}}(\boldsymbol{x}_i) = \sum_{i=1}^{N} \log \left(\sum_{z} p_{\boldsymbol{\theta}}(\boldsymbol{x}_i, z) \right)$$

... which is hard to optimize for all latent variable models (due to log of a sum)

Expectation-Maximization (EM)

Expectation-Maximization (EM) is a general algorithm for estimating latent variable models

- Most common application: Gaussian Mixture models
- ... but many other (deep) models as well
- Its extension is called Variational Bayes, which is underlying variational auto-encoder and other variational inference techniques
- Very hot research topic... pays off to look into the math of it

EM can be derived in 2 ways:

- Jensen's inequality (not covered)
- Decomposition in lower-bound and KL-term

Expectation-Maximization (EM)

EM uses a lower bound of the marginal log-likelihood for the optimization

• For simplicity, lets consider only a single data-point first



- Where q(z) is called the variational / auxiliary distribution
 - This decomposition holds for any q(z)
 - By introducing q(z), the optimization will become much simpler
- Why is that the same?
 - We can use Bayes rule for $p(z|x) = \frac{p(x, z|\theta)}{p(x|\theta)}$ and all terms except $p(x|\theta)$ cancel

Basics: Kullback-Leibler Divergences

The KL-divergence is a important similarity measure for distributions

$$\mathrm{KL}(q(\boldsymbol{x})||p(\boldsymbol{x})) = \sum_{\boldsymbol{x}} q(\boldsymbol{x}) \log \frac{q(\boldsymbol{x})}{p(\boldsymbol{x})}$$

- Its always non-negative
- If its zero, both distributions are the same:
- **It is non-symmetric** (hence, its not a distance metric):

 $KL(q||p) \ge 0$ $KL(q||p) = 0 \iff q = p$ $KL(q||p) \ne KL(p||q)$

- Can be used to find different approximations of distributions
- Used a lot in Variational Inference, Reinforcement Learning, Information theory...

EM-Decomposition

Derivation:

 $\log p(\boldsymbol{x}) = \sum \boldsymbol{q}(\boldsymbol{z}) \log p(\boldsymbol{x})$ 1. Introduce variational distribution q(z) $= \sum q(\boldsymbol{z}) \left(\log p(\boldsymbol{x}, \boldsymbol{z}) - \log p(\boldsymbol{z} | \boldsymbol{x}) \right)$ Use Bayesian theorem $p(x) = \frac{p(x, z)}{p(z|x)}$ 2. $= \sum q(z) \big(\log p(\boldsymbol{x}, z) - \log q(z) \big)$ 3. Add and subtract $\log q(z)$ $+\log q(z) - \log p(z|x)$ $= \sum_{z} q(z) \log \frac{p(\boldsymbol{x}, z)}{q(z)} + \sum_{z} q(z) \log \frac{q(z)}{p(z|\boldsymbol{x})}$ 4. Write as 2 sums Lower Bound $\mathcal{L}(q)$ KL Divergence: $KL(q(z)||p(z|\boldsymbol{x}))$ Gerhard Neumann | Machine Learning 1 | KIT | WS 2021/2022 24

EM Decomposition

Marginal Likelihood decomposes in 2 terms: $\log p(\boldsymbol{x}|\boldsymbol{\theta}) = \mathcal{L}(q, \boldsymbol{\theta}) + \mathrm{KL}(q(z)||p(z|\boldsymbol{x}))$

• Lower bound
$$\mathcal{L}(q, \theta) = \sum_{z} q(z) \log p(x, z | \theta) - \sum_{z} q(z) \log q(z)$$

Contains $\log p(\boldsymbol{x}, z | \boldsymbol{\theta})$ instead of $\log p(\boldsymbol{x} | \boldsymbol{\theta}) = \log \sum p(\boldsymbol{x}, z | \boldsymbol{\theta})$ _

- ... which is much easier to optimize (convex for most distributions) _
- Each $\log p(\boldsymbol{x}, z | \boldsymbol{\theta})$ is weighted by q(z)
- Why is it a lower bound? ٠
 - Since $\operatorname{KL}(q||p) \geq 0$ it follows that $\mathcal{L}(q, \theta) \leq \log p(\boldsymbol{x}|\boldsymbol{\theta})$

Expectation-Maximization Steps

EM iteratively applies 2 steps:

- (E)xpectation-step: $q(z) = \arg\min_{x} \operatorname{KL}(q(z)||p(z|x))$
 - Find q(z) that minimizes KL
 - Can be done in closed form for discrete z (e.g. mixtures):

$$q(z) = p(z|\boldsymbol{x}, \boldsymbol{\theta}_{\text{old}}) = \frac{p(\boldsymbol{x}, z | \boldsymbol{\theta}_{\text{old}})}{\sum_{z} p(\boldsymbol{x}, z | \boldsymbol{\theta}_{\text{old}})}$$

- Observations:
 - The marginal log-likelihood $\log p(\boldsymbol{x}|\boldsymbol{\theta})$ is unaffected by the E-step
 - As KL is minimized, lower bound has to go up
 - After the E-step KL(q(z)||p(z|x)) = 0 and therefore, the lower bound is tight, i.e.:

$$\log p(\boldsymbol{x}|\boldsymbol{\theta}) = \mathcal{L}(q, \boldsymbol{\theta})$$



Expectation-Maximization Steps

EM iteratively applies 2 steps:

• (M)aximization-step:

$$\boldsymbol{\theta} = \operatorname*{arg\,max}_{\boldsymbol{\theta}} \mathcal{L}(q, \boldsymbol{\theta}) = \operatorname*{arg\,max}_{\boldsymbol{\theta}} \sum_{z} q(z) \log p(\boldsymbol{x}, z | \boldsymbol{\theta}) + \mathrm{const}$$

- Maximize lower bound with respect to ${m heta}$
- Also called the complete-data likelihood
- Each possible value of the missing data is weighted by

$$q(z) = p(z|\boldsymbol{x}, \boldsymbol{\theta}_{\mathrm{old}})$$



EM Convergence Properties

EM improves the lower bound

 $\mathcal{L}(q_{\text{new}}, \boldsymbol{\theta}_{\text{new}}) \geq \mathcal{L}(q_{\text{old}}, \boldsymbol{\theta}_{\text{old}})$

- M-step: Lower bound is maximized
- E-step: KL is set to 0, lower bound has to go up
- EM improves the marginal likelihood

 $\log p(\boldsymbol{x}|\boldsymbol{\theta}_{\text{new}}) \geq \log p(\boldsymbol{x}|\boldsymbol{\theta}_{\text{old}})$

- M-step: Lower bound increases and KL increases (can't get smaller than 0)
- E-step: Marginal likelihood is unaffected





Illustration of EM

- Lower bound (blue curve) is a convex approximation of the marginal likelihood (red curve)
 - Maximum of lower bound can be easily obtained ($\theta^{
 m new}$)
 - Closed form solutions available, no gradient descent required
- Compute new lower bound for ${m heta}^{
 m new}$ (green curve)
- Due to the local approximation of the lowerbound, EM can only find local optima



EM for full dataset

For all data-points, the lower bound is given by:

$$\mathcal{L}(q, \boldsymbol{\theta}) = \sum_{i} \left(\int_{z} q_{i}(z) \log p(\boldsymbol{x}_{i}, z | \boldsymbol{\theta}) dz - \int_{z} q_{i}(z) \log q_{i}(z) dz \right)$$

- One latent variable z_i per data-point
- If z is discrete with K different values, than

 $q_i(z=k) = p(z=k|\boldsymbol{x}_i, \boldsymbol{\theta}_{\mathrm{old}})$

can be represented as a N x K matrix

• We will write $q_{ik} = q_i(z = k)$

Practical considerations...

How many mixture components do we need?

- More components will typically lead to a better likelihood
- But are more components necessarily better? Not always, because of overfitting!
- It's again a model-selection problem (cross-validate on a validation-set)
- Bayesian methods can be used to integrate out number of components (tricky to get them to work)

How do we initialize:

- EM can give very poor results with wrong initialization
- Most common approach:
 - Use k-means (simple clustering algorithm) to initialize the centers
 - Use a fixed value for the covariance

EM for Dimensionality Reduction

EM for Dimensionality Reduction (aka. probabilistic PCA)

We can also formulate dimensionality reduction using latent variables

Idea: Introduce a latent variable model to relate a D-dimensional observation vector to a corresponding M-dimensional gaussian latent variable (with M < D)

$$oldsymbol{x} = oldsymbol{W}oldsymbol{z} + oldsymbol{\mu} + oldsymbol{\epsilon}$$

- **z** is a d' latent variable (our low dimensional representation)
- **W** is a *D* x *M* matrix relating the latent space **z** with the original space **x**
- μ is a constant offset vector
- $\boldsymbol{\epsilon}$ is a d-dimensional Gaussian noise vector $\boldsymbol{\epsilon} \sim \mathcal{N}(\boldsymbol{0}, \sigma^2 \boldsymbol{I})$



EM for Dimensionality Reduction

Probabilistic Dimensionality Reduction Model:

- $\boldsymbol{z} \in \mathbb{R}^M, \quad \boldsymbol{x} \in \mathbb{R}^D, \quad M < D$
- Continuous Latent Variable:
 - $p(\boldsymbol{z}) = \mathcal{N}(\boldsymbol{0}, \boldsymbol{I})$
 - Assume 0 mean, unit variance distribution in latent space
- Observation Model

 $p(\boldsymbol{x}|\boldsymbol{z}, \boldsymbol{\theta}) = \mathcal{N}(\boldsymbol{W}\boldsymbol{z} + \boldsymbol{\mu}, \sigma^2 \boldsymbol{I})$

– with parameters $oldsymbol{ heta} = \{oldsymbol{W}, oldsymbol{\mu}, \sigma^2\}$



Generative Process

Our model can be interpreted in terms of a generative process

1. Sample latent variable

 $oldsymbol{z} \sim \mathcal{N}(oldsymbol{0}, oldsymbol{I})$

- 2. Linearly project to high-D space
 - $oldsymbol{y} = oldsymbol{W}oldsymbol{z} + oldsymbol{\mu}$
- 3. Sample noise

 $oldsymbol{\epsilon} \sim \mathcal{N}(oldsymbol{0}, \sigma^2 oldsymbol{I})$

4. Add noise to obtain **x**

 $x = y + \epsilon$



Marginal likelihood

Marginal likelihood is given by:

$$p(\boldsymbol{x}|\boldsymbol{\theta}) = \int_{\boldsymbol{z}} p(\boldsymbol{x}|\boldsymbol{z}, \boldsymbol{\theta}) p(\boldsymbol{z}) d\boldsymbol{z} = \int_{\boldsymbol{z}} \mathcal{N}(\boldsymbol{x}|\boldsymbol{W}\boldsymbol{z} + \boldsymbol{\mu}, \sigma^2 \boldsymbol{I}) \mathcal{N}(\boldsymbol{z}|\boldsymbol{0}, \boldsymbol{I}) d\boldsymbol{z}$$

Maximize the marginal log-likelihood:

$$\operatorname{loglike}(\boldsymbol{\theta}) = \sum_{i=1}^{N} \log p(\boldsymbol{x}|\boldsymbol{\theta}) = \sum_{i=1}^{N} \log \left(\int_{\boldsymbol{z}} \mathcal{N}(\boldsymbol{x}|\boldsymbol{W}\boldsymbol{z} + \boldsymbol{\mu}, \sigma^{2}\boldsymbol{I}) \mathcal{N}(\boldsymbol{z}|\boldsymbol{0}, \boldsymbol{I}) d\boldsymbol{z} \right)$$

This is a typical case for using EM

- It can however also be solved in closed form as everything is Gaussian and linear
- But it is somehow complex and using EM is a much more general solution
- Its a good example to understand EM

Expectation-step

We need to compute the posterior distribution



• Application of Bayes' Rule with Gaussian distributions

$$p(\boldsymbol{x}|\boldsymbol{z}, \boldsymbol{\theta}) = \mathcal{N}(\boldsymbol{W}\boldsymbol{x} + \boldsymbol{\mu}, \sigma^2 \boldsymbol{I}), \quad p(\boldsymbol{z}) = \mathcal{N}(\boldsymbol{0}, \boldsymbol{I})$$

• Posterior is Gaussian with mean and variance

$$\boldsymbol{\mu}_{\boldsymbol{z}|\boldsymbol{x}_i} = (\boldsymbol{W}^T \boldsymbol{W} + \sigma^2 \boldsymbol{I})^{-1} \boldsymbol{W}^T (\boldsymbol{x}_i - \boldsymbol{\mu}), \qquad \boldsymbol{\Sigma}_{\boldsymbol{z}|\boldsymbol{x}_i} = \sigma^2 (\boldsymbol{W}^T \boldsymbol{W} + \sigma^2 \boldsymbol{I})^{-1}$$

- Not covered now, see Lecture 9, Bayesian Learning
- Only the case because x is linear in z!

Maximization Step

Maximize the lower bound with respect to θ ...

$$\begin{aligned} \mathcal{L}(q,\boldsymbol{\theta}) &= \sum_{i} \left(\int_{\boldsymbol{z}} q_{i}(\boldsymbol{z}) \log p(\boldsymbol{x}_{i}, \boldsymbol{z} | \boldsymbol{\theta}) d\boldsymbol{z} - \int_{\boldsymbol{z}} q_{i}(\boldsymbol{z}) \log q_{i}(\boldsymbol{z}) d\boldsymbol{z} \right) \\ &= \sum_{i} \left(\int_{\boldsymbol{z}} q_{i}(\boldsymbol{z}) \left(\log p(\boldsymbol{x}_{i} | \boldsymbol{z}, \boldsymbol{\theta}) + \log p(\boldsymbol{z}) \right) d\boldsymbol{z} - \int_{\boldsymbol{z}} q_{i}(\boldsymbol{z}) \log q_{i}(\boldsymbol{z}) d\boldsymbol{z} \right) \\ &= \sum_{i} \int_{\boldsymbol{z}} q_{i}(\boldsymbol{z}) \log p(\boldsymbol{x}_{i} | \boldsymbol{z}, \boldsymbol{\theta}) d\boldsymbol{z} + \underbrace{\operatorname{const}}_{\text{independent of } \boldsymbol{\theta}} = \sum_{i} \mathbb{E}_{q_{i}(\boldsymbol{z})} \left[\log p(\boldsymbol{x}_{i} | \boldsymbol{z}, \boldsymbol{\theta}) \right] + \underbrace{\operatorname{const}}_{\text{independent of } \boldsymbol{\theta}} \end{aligned}$$

- Continuous latent variable: How can we solve the integral?
 - a) $q_i(z)$ is Gaussian, can be solved in principle in closed form (not covered)
 - b) Simpler: Sampling! I.e, we can use Monte Carlo Estimates

Recap: Monte-carlo estimation (Lecture 2)

Expectations can always be approximated by samples:

$$\mathbb{E}_p[f(x)] = \int p(x)f(x)dx \approx \frac{1}{N} \sum_{x_i \sim p(x)} f(x_i)$$

• Necessary if no analytical solution exists to compute the integral (typical case)

Maximization Step

Monte-Carlo estimate for the lower bound

$$\mathcal{L}(q, \boldsymbol{\theta}) = \sum_{i} \mathbb{E}_{q_i(\boldsymbol{z})} \left[\log p(\boldsymbol{x}_i | \boldsymbol{z}, \boldsymbol{\theta}) \right] \approx \sum_{i} \frac{1}{N} \sum_{z_{ik} \sim q_i(\boldsymbol{z})} \log p(\boldsymbol{x}_i | \boldsymbol{z}_{ik}, \boldsymbol{\theta})$$

• If we only use a single sample z_i per i (i.e. N = 1), we get

$$\mathcal{L}(q, oldsymbol{ heta}) pprox \sum_i \log p(oldsymbol{x}_i | oldsymbol{z}_i, oldsymbol{ heta}), ext{ where } oldsymbol{z}_i \sim q_i(oldsymbol{z}).$$

- Maximizing $\mathcal{L}(q, \theta)$ is a standard maximum likelihood problem with Gaussian linear models.
- We know the solution already (standard least squares):

$$\begin{bmatrix} \boldsymbol{\mu} \\ \boldsymbol{W} \end{bmatrix} = (\boldsymbol{Z}^T \boldsymbol{Z})^{-1} \boldsymbol{Z}^T \boldsymbol{X}, \text{ with } \boldsymbol{Z} = \begin{bmatrix} 1 & \boldsymbol{z}_1^T \\ \vdots & \vdots \\ 1 & \boldsymbol{z}_n^T \end{bmatrix} \text{ and } \boldsymbol{X} = \begin{bmatrix} \boldsymbol{x}_1^T \\ \vdots \\ \boldsymbol{x}_n^T \end{bmatrix}$$
$$\sigma^2 = \frac{1}{nd} \sum_{i=1}^n \sum_{k=1}^d (y_{ik} - x_{ik})^2, \text{ with } \boldsymbol{y}_i = \boldsymbol{W} \boldsymbol{z}_i + \boldsymbol{\mu}$$

Algorithm: EM for PCA

- Initialize: Use average of \mathbf{x} for $\boldsymbol{\mu}$, random matrix \mathbf{W}
- **Repeat** until convergence:
 - Expectation-step:
 - Compute posterior mean and covariance

$$\boldsymbol{\mu}_{\boldsymbol{z}|\boldsymbol{x}_i} = (\boldsymbol{W}^T \boldsymbol{W} + \sigma^2 \boldsymbol{I})^{-1} \boldsymbol{W}^T (\boldsymbol{x}_i - \boldsymbol{\mu}), \qquad \boldsymbol{\Sigma}_{\boldsymbol{z}|\boldsymbol{x}_i} = \sigma^2 (\boldsymbol{W}^T \boldsymbol{W} + \sigma^2 \boldsymbol{I})^{-1}$$

• Generate latent samples:

$$oldsymbol{z}_i \sim \mathcal{N}(oldsymbol{\mu}_{oldsymbol{z} \mid oldsymbol{x}_i}, oldsymbol{\Sigma}_{oldsymbol{z} \mid oldsymbol{x}_i})$$

- Maximization-step: Update $\boldsymbol{W}, \ \boldsymbol{\mu} \ \mathrm{and} \ \sigma^2$

$$\begin{bmatrix} \boldsymbol{\mu} \\ \boldsymbol{W} \end{bmatrix} = (\boldsymbol{Z}^T \boldsymbol{Z})^{-1} \boldsymbol{Z}^T \boldsymbol{X}, \qquad \sigma^2 = \frac{1}{nd} \sum_{i=1}^n \sum_{k=1}^d (y_{ik} - x_{ik})^2$$

Illustration: EM for PCA



Probabilistic PCA vs. PCA

Typically, PCA using eigenvector decomposition is preferred:

- Single one-step solution
- Very fast

However, looking at EM for dimensionality reduction makes sense if:

- We need a density
- Helps us to understand EM
- Helps us to understand more complex dimensionality reduction methods (variational auto encoders use the same principles)

Additional Notes for EM

EM assumes that E-step can set the KL to zero:

- I.e. we can evaluate the posterior analytically
- Lower bound is tight
- Marginal likelihood always improves (good to check for debugging!)
- Only possible if z is discrete or we have linear Gaussian models!

For more complex latent variable models (e.g. Deep Neural Networks), this is typically not possible:

- Extension of EM called Variational Bayes / Variational Inference can still do that
- Approximates the posterior, i.e. KL will be > 0 after E-step
- Very active research, underlying algorithm of many deep learning architectures (e.g. variational autoencoder)
- Will be covered in the end of the lecture

Takeaway messages

You know now:

- The difference between parametric and non-parametric models
- Different non-parametric models (histogram, kernel density estimation and k-nearest neighbors)
- What mixture models and latent variable models are
- What the Expectation-Maximization idea and algorithm are
- Why does EM converge
- How to apply EM to GMMs (discrete latent variables) and PCA (continuous latent variables)



Self-test questions

- What are mixture models?
- Should gradient methods be used for training mixture models?
- How does the EM algorithm work?
- What is the biggest problem of mixture models?
- How does EM decomposes the marginal likelihood?
- Why does EM always improve the lower bound?
- Why does EM always improve the marginal likelihood?
- Why can we optimize each mixture component independently with EM
- Why do we need sampling for continuous latent variables?