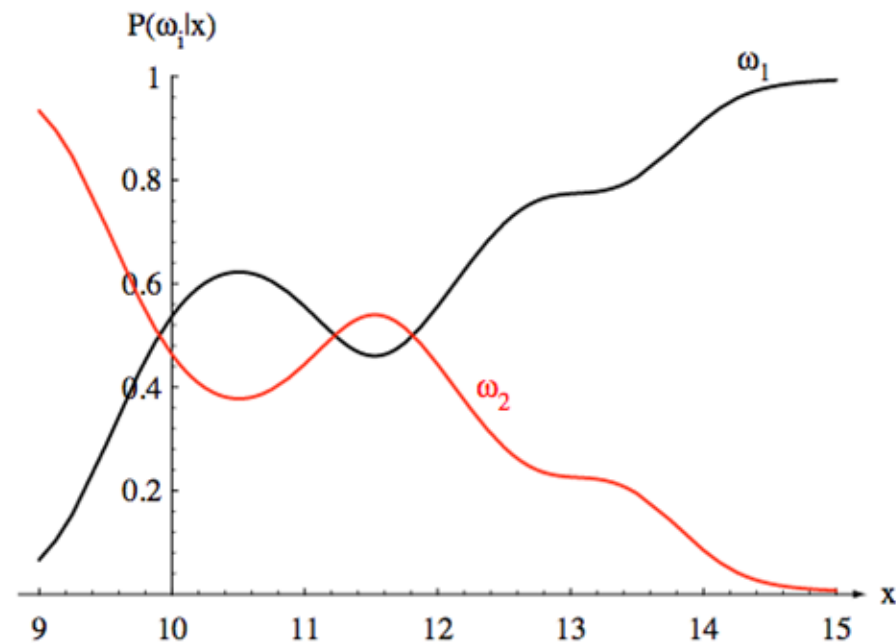
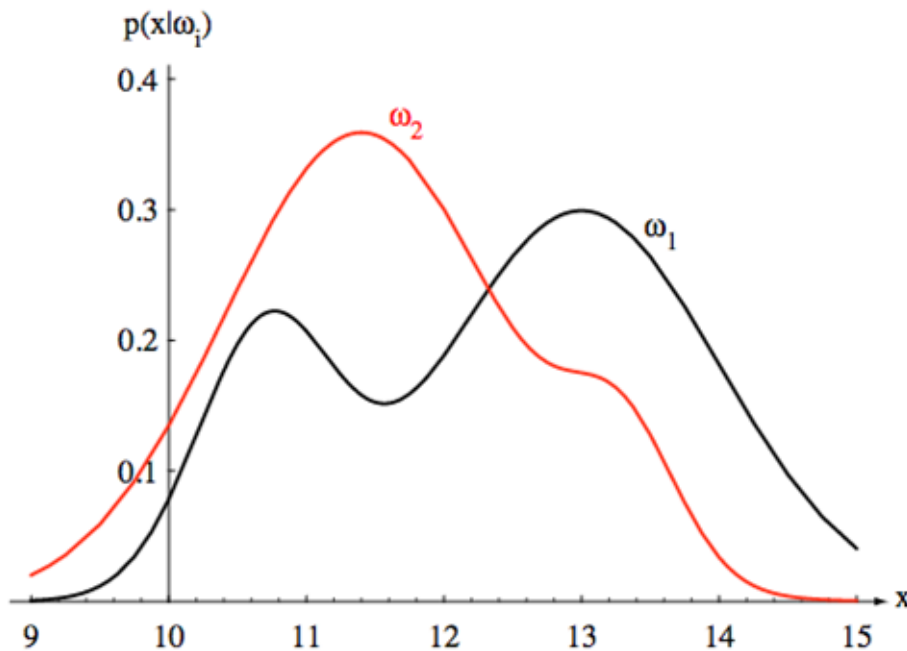


Recap

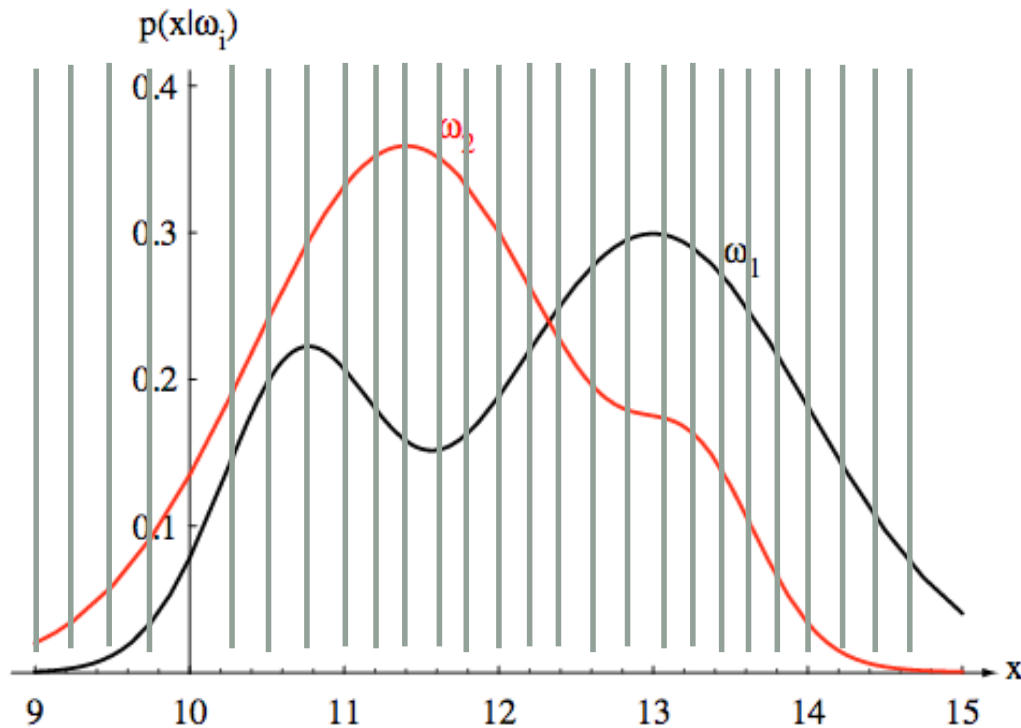
A simple decision rule

- If we can know either $p(x|w)$ or $p(w|x)$ we can make a classification guess



Goal: Find $p(x|w)$ or $p(w|x)$

A simple way to estimate $p(x|w)$



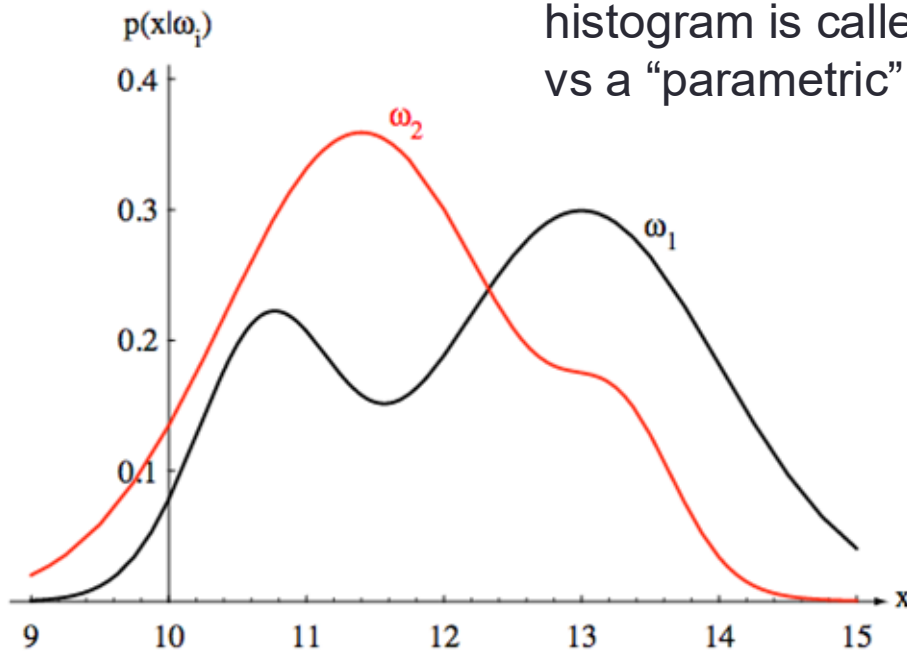
Make a histogram!

What happens if there is no data in a histogram bin?

The parametric approach

- We **assume** $p(x|w)$ or $p(w|x)$ follow some distributions with parameter θ

The method where we model the distribution using a histogram is called a “non-parametric” approach vs a “parametric” approach



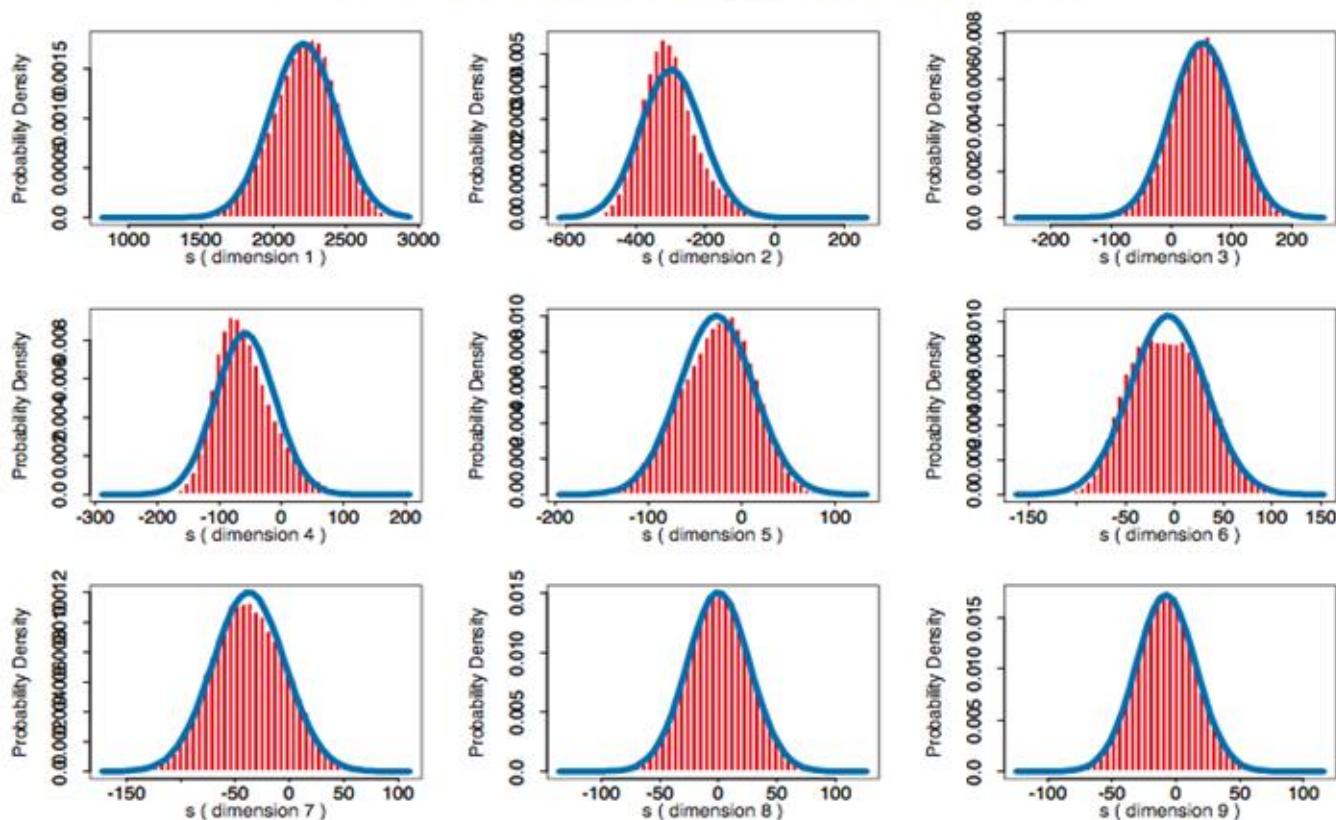
$$p(x) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-(x-\mu)^2/2\sigma^2}$$

Goal: Find θ so that we can estimate $p(x|w)$ or $p(w|x)$

GMM

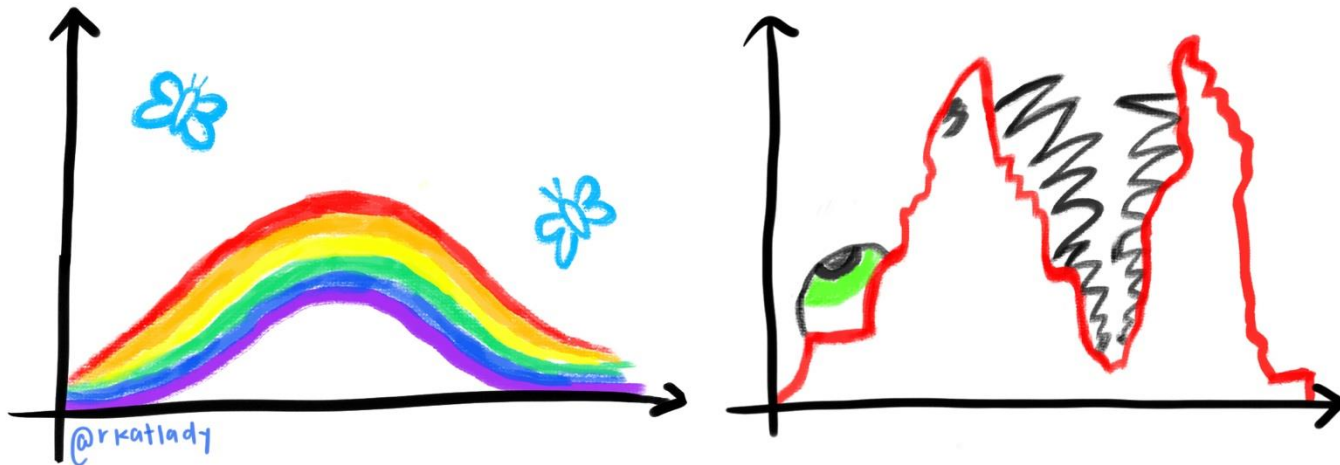
Can we do better than a single Gaussian?

First 9 MFCC's from [s]: Gaussian PDF



UNDERLYING DISTRIBUTIONS:

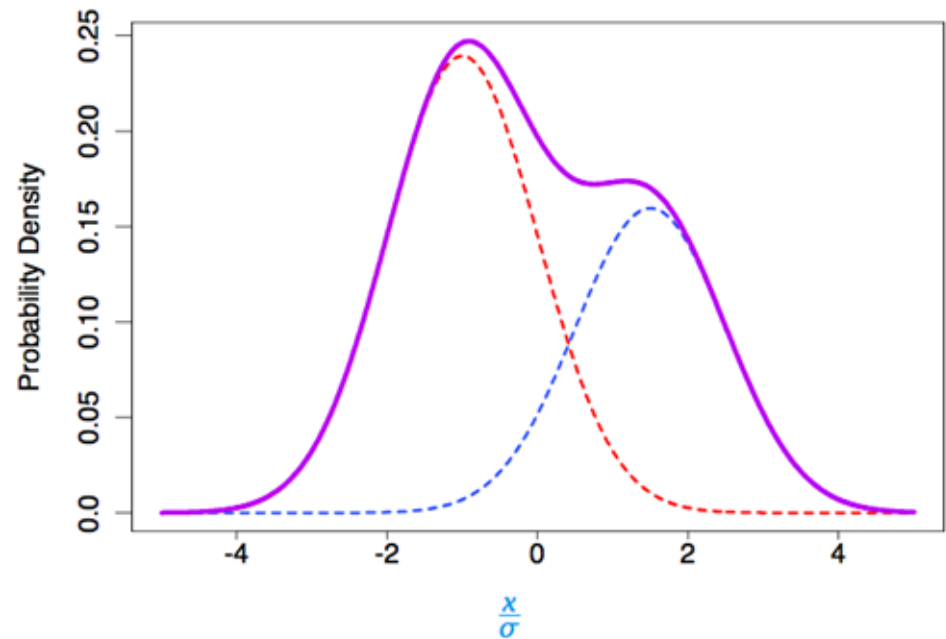
PARAMETRIC
ASSUMPTIONS VS. REALITY



Gaussian Mixture Models (GMMs)

- Gaussians cannot handle multi-modal data well
- Consider a class can be further divided into additional factors
- Mixing weight makes sure the overall probability sums to 1

$$P(x) \sim \sum_{k=1}^K w_k N(\mu_k, \sigma_k)$$

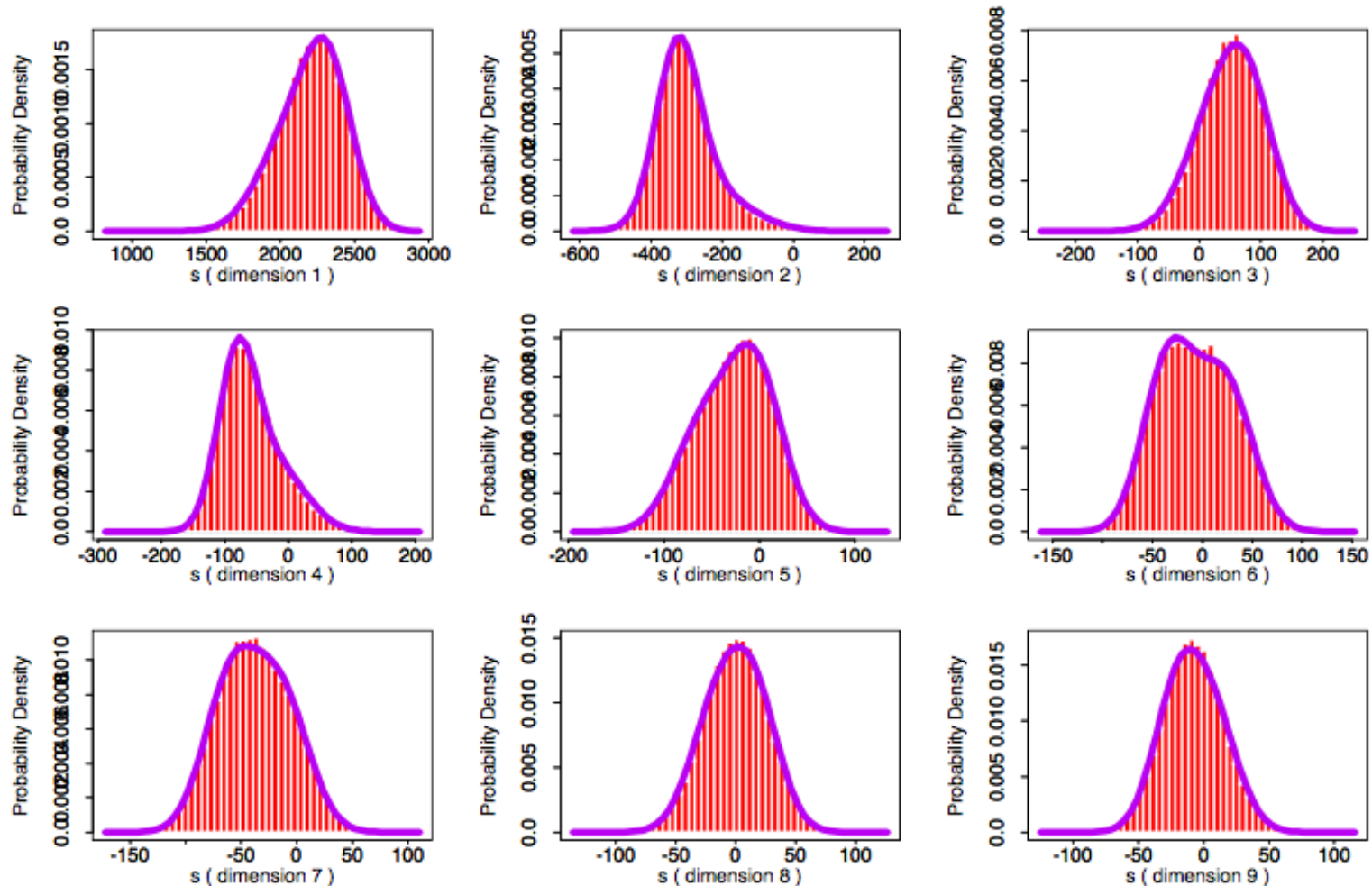


$$p(x) = 0.6p_1(x) + 0.4p_2(x)$$

$$p_1(x) \sim N(-\sigma, \sigma^2) \quad p_2(x) \sim N(1.5\sigma, \sigma^2)$$

Mixture of two Gaussians

[s]: 2 Gaussian Mixture Components/Dimension



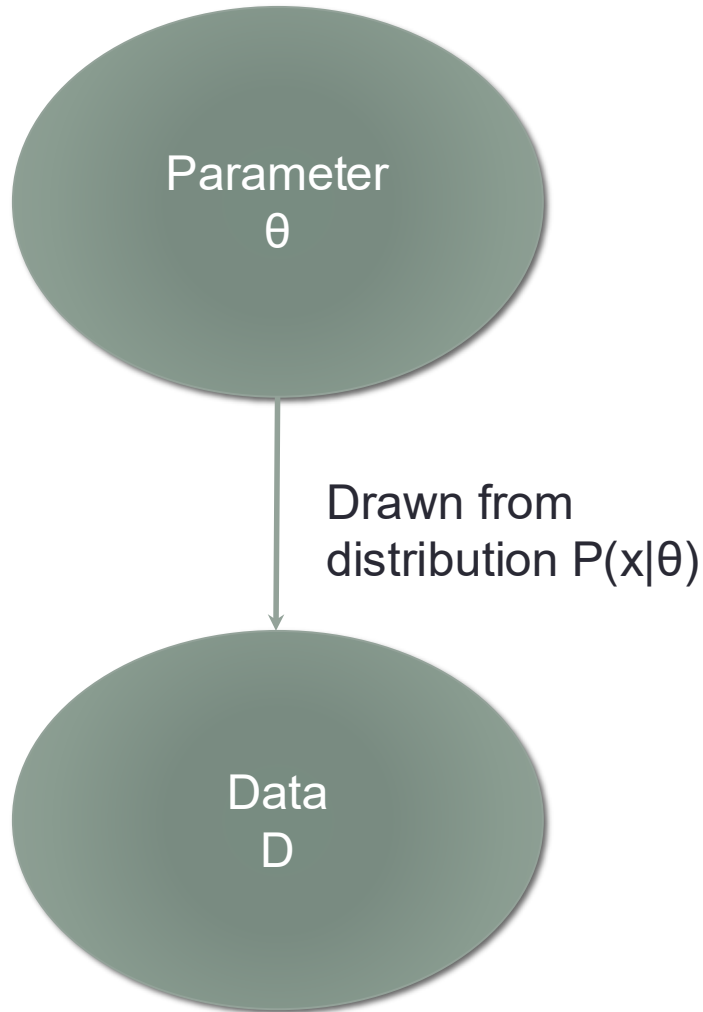
Mixture models

$$p(x) = \sum_k p(k)p_k(x)$$

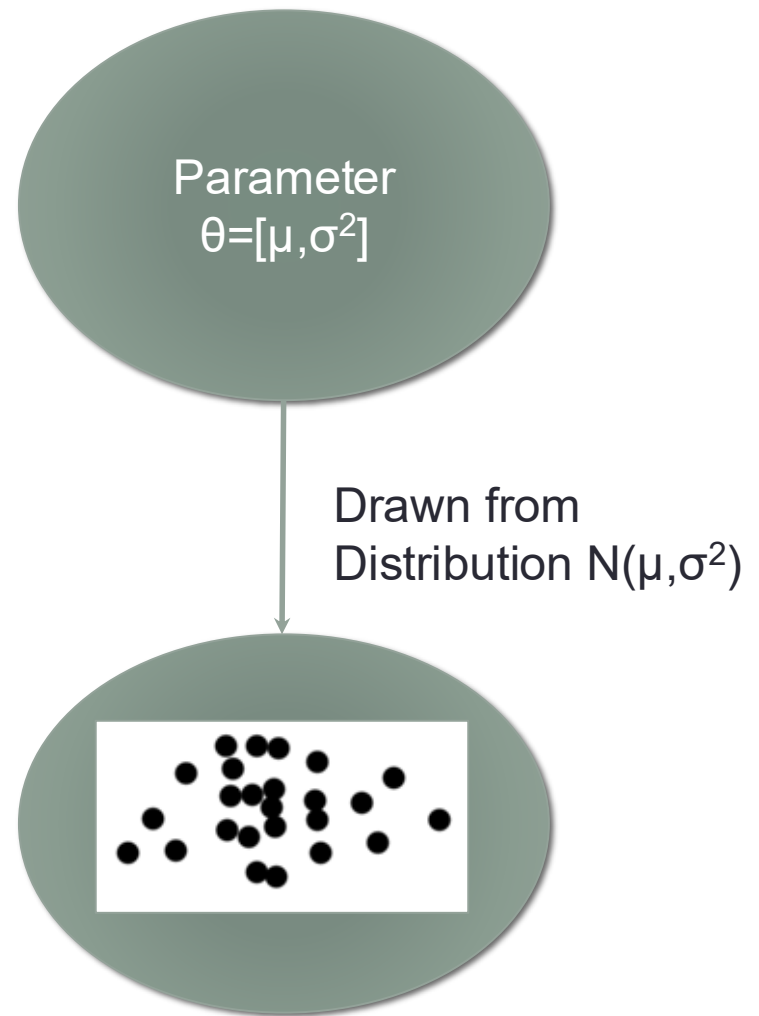
- A mixture of models from the same distributions (but with different parameters)
- Different mixtures can come from different sub-class
 - Cat class
 - Siamese cats
 - Persian cats
- $p(k)$ is usually categorical (discrete classes)
- Usually the exact class for a sample point is unknown.
 - Latent variable

Parametric models

Parametric models

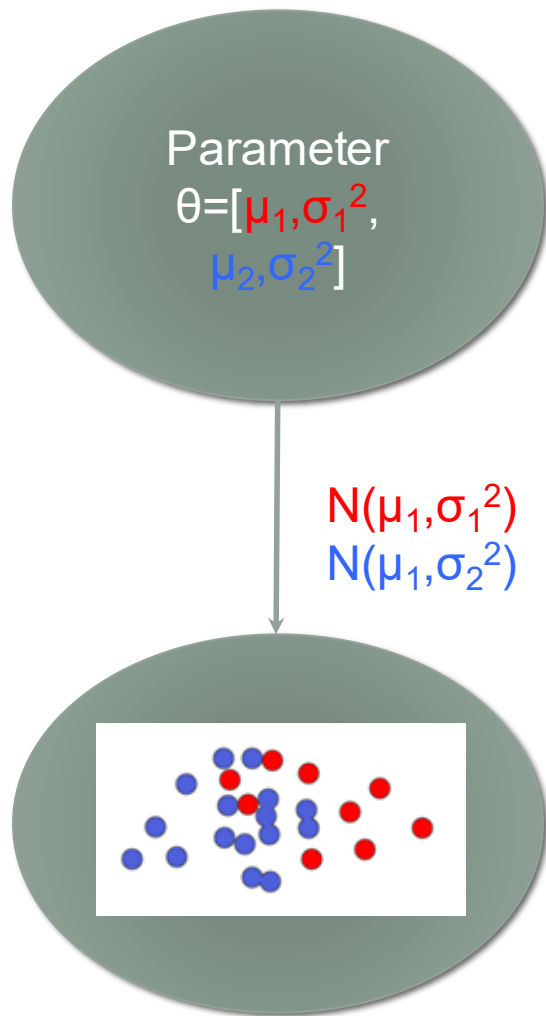


Gaussian

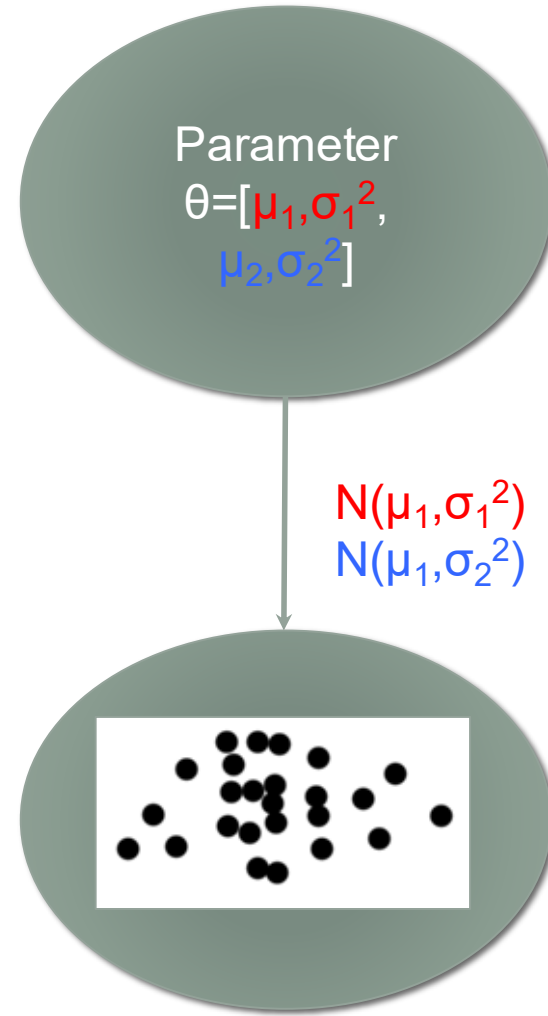


What if some data is missing?

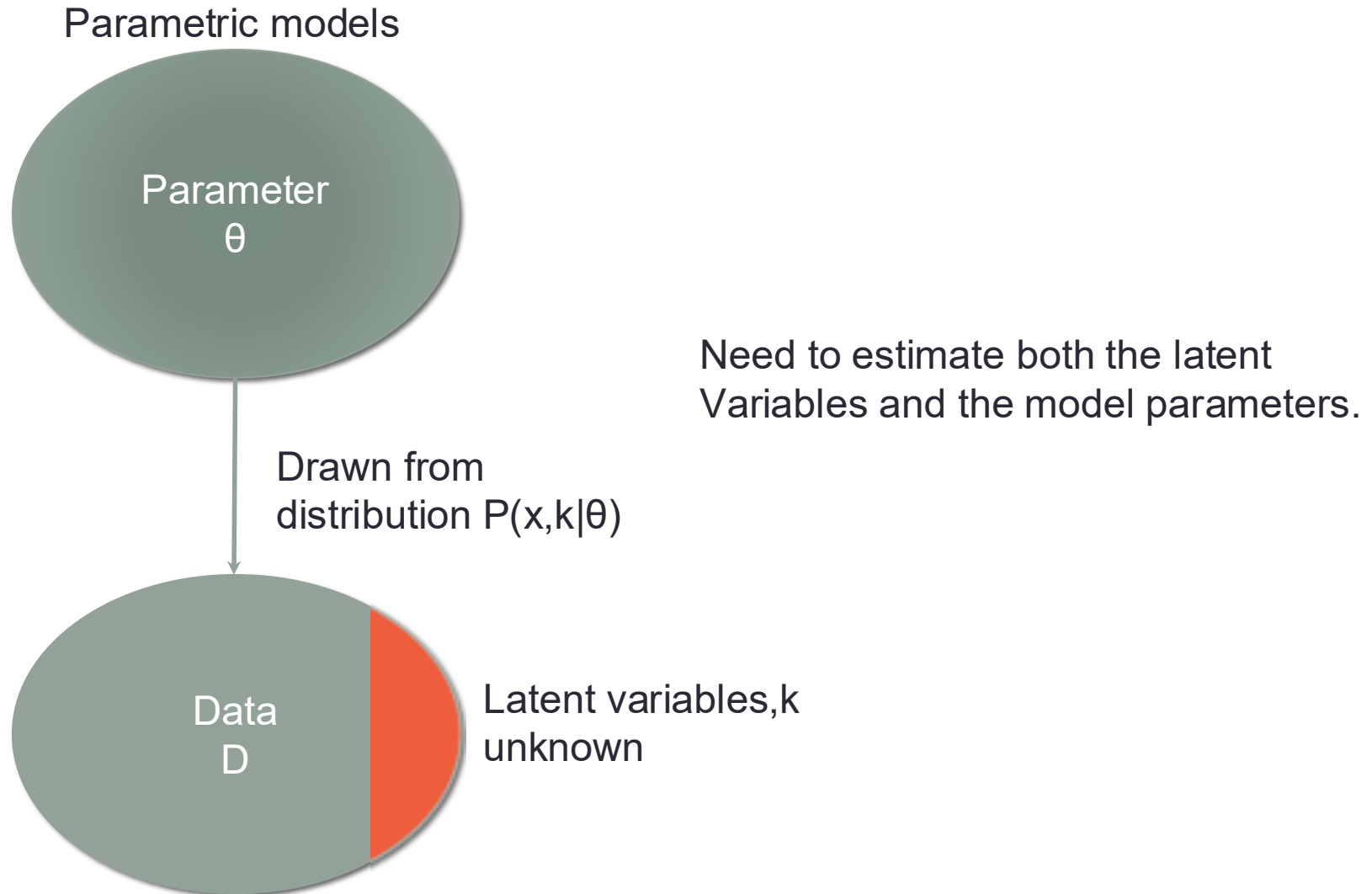
Mixture of Gaussian



Unknown mixture labels



Estimating missing data



Slight difference in notation

$p(\mathbf{x}|\theta)$ vs $p(\mathbf{x};\theta)$

θ as a RV at a fixed value vs θ as a fixed parameter

Most of the time can be used interchangeably

Estimating latent variables and model parameters

- GMM $p(x) = \sum_k p(k)N(\mu_k, \sigma_k)$
- Observed (x_1, x_2, \dots, x_N)
- Latent (k_1, k_2, \dots, k_N) from K possible mixtures
- Parameter for $p(k)$ is ϕ , $p(k = 1) = \phi_1$, $p(k = 2) = \phi_2 \dots$

$$l(\phi, \mu, \Sigma) = \sum_{n=1}^N \log p(x^{(i)}; \phi, \mu, \sigma)$$

$$= \sum_{n=1}^N \log \sum_{l=1}^K p(x_n | k_{n,l}; \mu, \sigma) p(k_{n,l}; \phi)$$

Make things hard to solve

Cannot be solved by differentiating

Assuming k

- What if we somehow know k_n ?
- Maximizing wrt to ϕ , μ , σ gives

$$\phi_j = \frac{1}{N} \sum_{n=1}^N 1(k_n = j)$$

$$\mu_j = \frac{\sum_{n=1}^N 1(k_n = j) x_n}{\sum_{n=1}^N 1(k_n = j)}$$

$$\sigma_j^2 = \frac{\sum_{n=1}^N 1(k_n = j) (x_n - \mu_j)^2}{\sum_{n=1}^N 1(k_n = j)}$$

$1(\text{condition})$

Indicator function. Equals one if condition is met. Zero otherwise

Iterative algorithm

- Initialize ϕ , μ , σ
- Repeat till convergence
 - Expectation step (E-step) : Estimate the latent labels \mathbf{k}
 - Maximization step (M-step) : Estimate the parameters ϕ , μ , σ given the latent labels
- Called Expectation Maximization (EM) Algorithm
- How to estimate the latent labels?

Iterative algorithm

- Initialize ϕ, μ, σ
- Repeat till convergence
 - **Expectation step** (E-step) : Estimate the latent labels \mathbf{k} by finding the pdf of k given everything else $p(k | x; \phi, \mu, \sigma)$
 - **Maximization step** (M-step) : Estimate the parameters ϕ, μ, σ given the latent labels by maximizing the **expectation** of the **log likelihood**
- Extension of MLE for latent variables
 - MLE : $\operatorname{argmax} \log p(x; \theta)$
 - EM : $\operatorname{argmax} \log \sum_k p(x, k; \theta)$

How to evaluate $\log \sum_k p(x, k; \theta)$ when we don't know k ?

Convex functions and Jensen's inequality

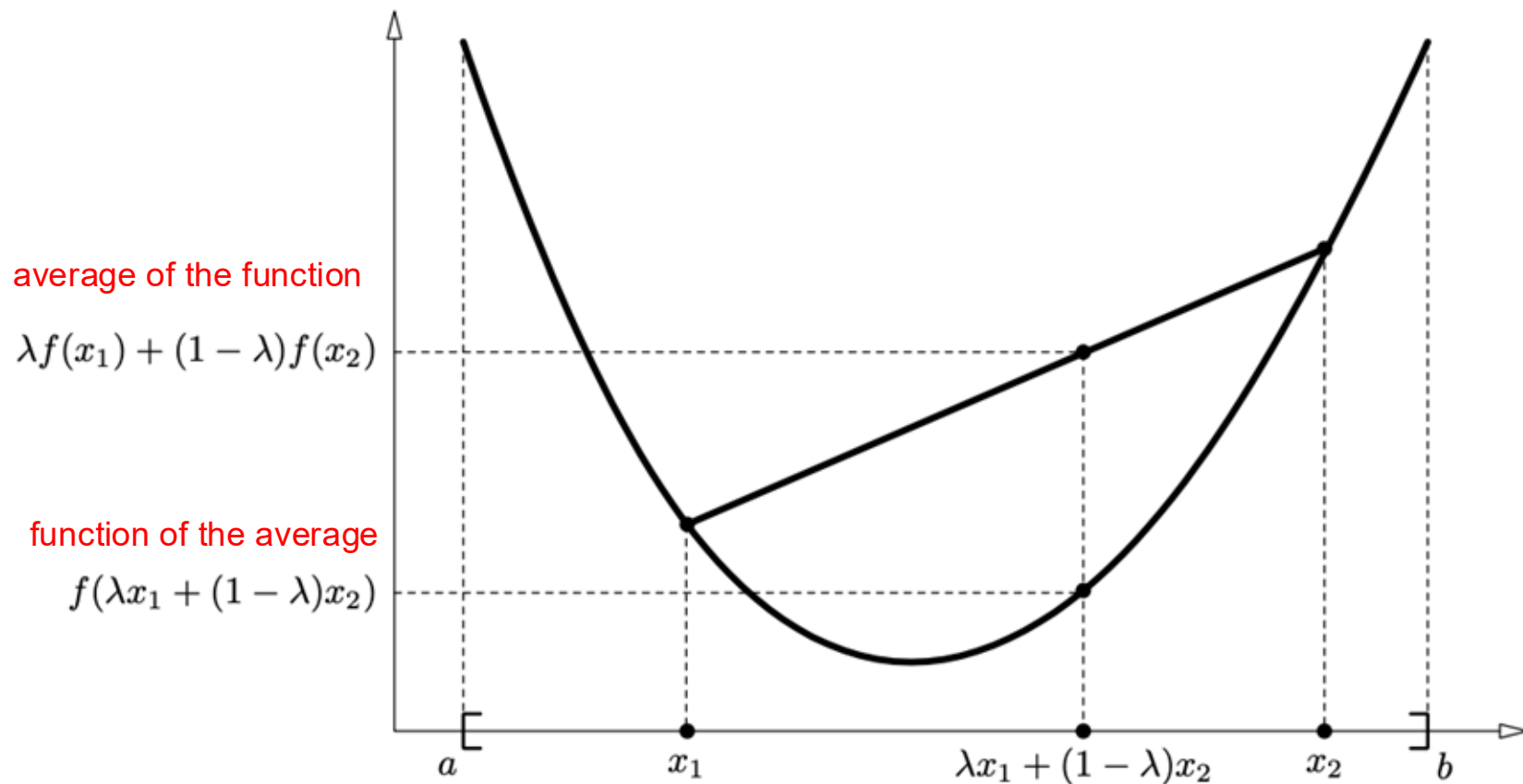


Figure 1: f is *convex* on $[a, b]$ if $f(\lambda x_1 + (1 - \lambda)x_2) \leq \lambda f(x_1) + (1 - \lambda)f(x_2)$
 $\forall x_1, x_2 \in [a, b], \lambda \in [0, 1]$.

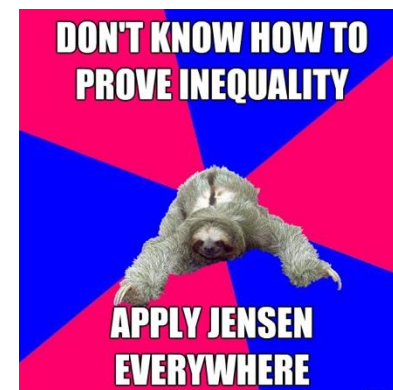
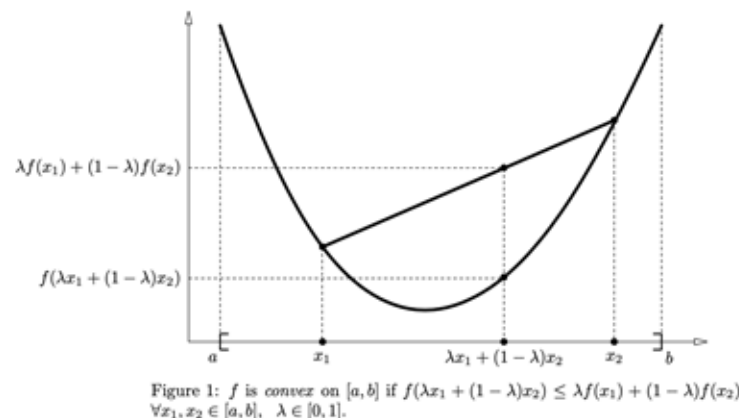
Jensen's inequality

Let f be a convex function on interval I

If x_1, x_2, \dots, x_n is in I ,

$w_1, \dots, w_n > 0$ and sums to 1
then,

$$f\left(\sum_i^n w_i x_i\right) \leq \sum_i^n w_i f(x_i)$$



If f is concave, flip the inequality.

Can view this as expectation

$$f(E[X]) \leq E[f(X)]$$

Jensen's inequality and ELBO

$\log \sum_k p(x, k; \theta)$

$$f\left(\sum_i^n w_i x_i\right) \leq \sum_i^n w_i f(x_i)$$

Maximize Evidence Lower Bound (ELBO) = $\sum_k Q(k) \log (p(x, k; \theta) / Q(k))$

Making the lower bound tight

We will make the bound tight for fixed θ
 Jensen's inequality is tight when?

$$f\left(\sum_i^n w_i x_i\right) \leq \sum_i^n w_i f(x_i)$$

$$f(E[X]) \leq E[f(X)]$$

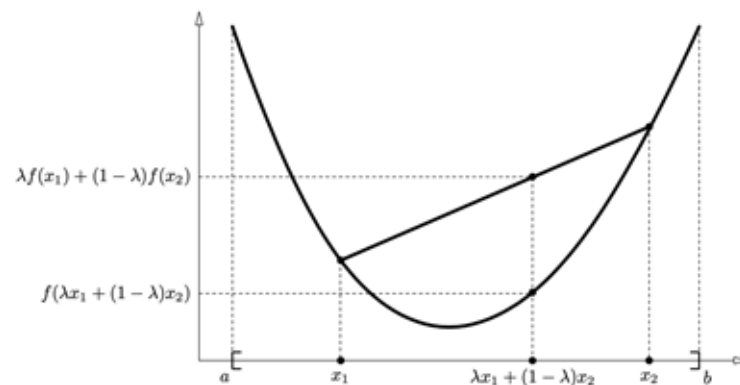


Figure 1: f is convex on $[a, b]$ if $f(\lambda x_1 + (1 - \lambda)x_2) \leq \lambda f(x_1) + (1 - \lambda)f(x_2)$
 $\forall x_1, x_2 \in [a, b], \lambda \in [0, 1]$.

Making the lower bound tight

We will make the bound tight for a fixed θ

$$f\left(\sum_i^n w_i x_i\right) \leq \sum_i^n w_i f(x_i)$$

$$f(E[X]) \leq E[f(X)]$$

If $f(\cdot)$ is strictly convex, Jensen's inequality is tight IFF

x_i are all equal

$E[X] = X = \text{constant}$

Making the lower bound tight

We will make the bound tight for a fixed θ

Jensen's inequality is tight when the inside of the expectation is a constant, c wrt the expectation of k

$$p(x, k; \theta) / Q(k) = c$$

or $Q(k) = p(k \mid x; \theta)$

Iterative algorithm (general)

- Goal of EM : $\log \sum_k p(x, k; \theta) \geq \sum_k Q(k) \log (p(x, k; \theta)/Q(k))$
- Maximize the ELBO instead
- Initialize Θ
- Repeat till convergence
 - Expectation step (E-step) : estimate the conditional expectation (the posterior)
 $Q(k) = p(k|x; \theta)$ using the current θ .
 - Maximization step (M-step) : Estimate new Θ given by maximizing the **ELBO** given current $Q(k)$

EM on a simple example

- Grades in class $P(A) = 0.5$ $P(B) = 0.5 - \theta$ $P(C) = \theta$
- We want to estimate θ from three known numbers
 - N_a N_b N_c
- Find the maximum likelihood estimate of θ

EM on a simple example

- Grades in class $P(A) = 0.5$ $P(B) = 0.5 - \theta$ $P(C) = \theta$
- We want to estimate θ from ONE known number
 - N_c (we also know N the total number of students)
- Find θ using EM

Will this work?

For iteration i , with $\theta^{(i)}$

$$\log \sum_k p(x, k; \theta^{(i)}) \geq \underbrace{\sum_k Q(k) \log (p(x, k; \theta^{(i)})/Q(k))}_{\text{ELBO}}$$

E-step, making the bound tight by picking $Q'(k)$ yields

$$\log \sum_k p(x, k; \theta^{(i)}) = \sum_k Q'(k) \log (p(x, k; \theta^{(i)})/Q'(k))$$

M-step, maximize ELBO by finding $\theta^{(i+1)}$

$$\sum_k Q'(k) \log (p(x, k; \theta^{(i)})/Q'(k)) \leq \sum_k Q'(k) \log (p(x, k; \theta^{(i+1)})/Q'(k))$$

For iteration $i+1$, with $\theta^{(i+1)}$

$$\log \sum_k p(x, k; \theta^{(i+1)}) \geq \sum_k Q(k) \log (p(x, k; \theta^{(i+1)})/Q(k))$$

Thus,

$$\log \sum_k p(x, k; \theta^{(i+1)}) \geq \log \sum_k p(x, k; \theta^{(i)})$$

So EM improves the likelihood at every step!

Notes on ELBO

We set $Q(k) = p(k \mid x; \theta)$ to make the inequality tight.

What if we cannot compute $p(k \mid x; \theta)$?

Use a looser bound by picking any $Q(k)$

Estimate $p(k \mid x; \theta)$ with $q(k \mid x; \theta)$ that we can compute

This is called **Variational Inference**

We will revisit this.

Estimating latent variables and model parameters

- GMM $p(x) = \sum_k p(k)N(\mu_k, \sigma_k)$
- Observed (x_1, x_2, \dots, x_N)
- Latent (k_1, k_2, \dots, k_N) from K possible mixtures
- Parameter for $p(k)$ is ϕ , $p(k = 1) = \phi_1$, $p(k = 2) = \phi_2 \dots$

$$l(\phi, \mu, \Sigma) = \sum_{n=1}^N \log p(x^{(i)}; \phi, \mu, \sigma)$$

$$= \sum_{n=1}^N \log \sum_{l=1}^K p(x_n | k_{n,l}; \mu, \sigma) p(k_{n,l}; \phi)$$

Make things hard to solve

Cannot be solved by differentiating

EM on GMM

- E-step

- Set soft labels: $w_{n,j}$ = probability that n th sample comes from j th mixture p

- Using Bayes rule

- $$p(k|x; \mu, \sigma, \phi) = \frac{p(x|k; \mu, \sigma, \phi) p(k; \mu, \sigma, \phi)}{p(x; \mu, \sigma, \phi)}$$

- $p(k|x; \mu, \sigma, \phi)$ is proportional to $p(x|k; \mu, \sigma, \phi) p(k; \phi)$

$$p(k_n = j | x_n; \phi, \mu, \Sigma) = \frac{N(\mu_j, \sigma_j) \phi_j}{\sum_l p(x_n; \mu_l, \sigma_l) p(k_n = l; \phi)}$$

EM on GMM

- M-step (hard labels)

$$\phi_j = \frac{1}{N} \sum_{n=1}^N 1(k_n = j)$$

$$\mu_j = \frac{\sum_{n=1}^N 1(k_n = j) x_n}{\sum_{n=1}^N 1(k_n = j)}$$

$$\sigma_j^2 = \frac{\sum_{n=1}^N 1(k_n = j) (x_n - \mu_j)^2}{\sum_{n=1}^N 1(k_n = j)}$$

EM on GMM

- M-step (soft labels)

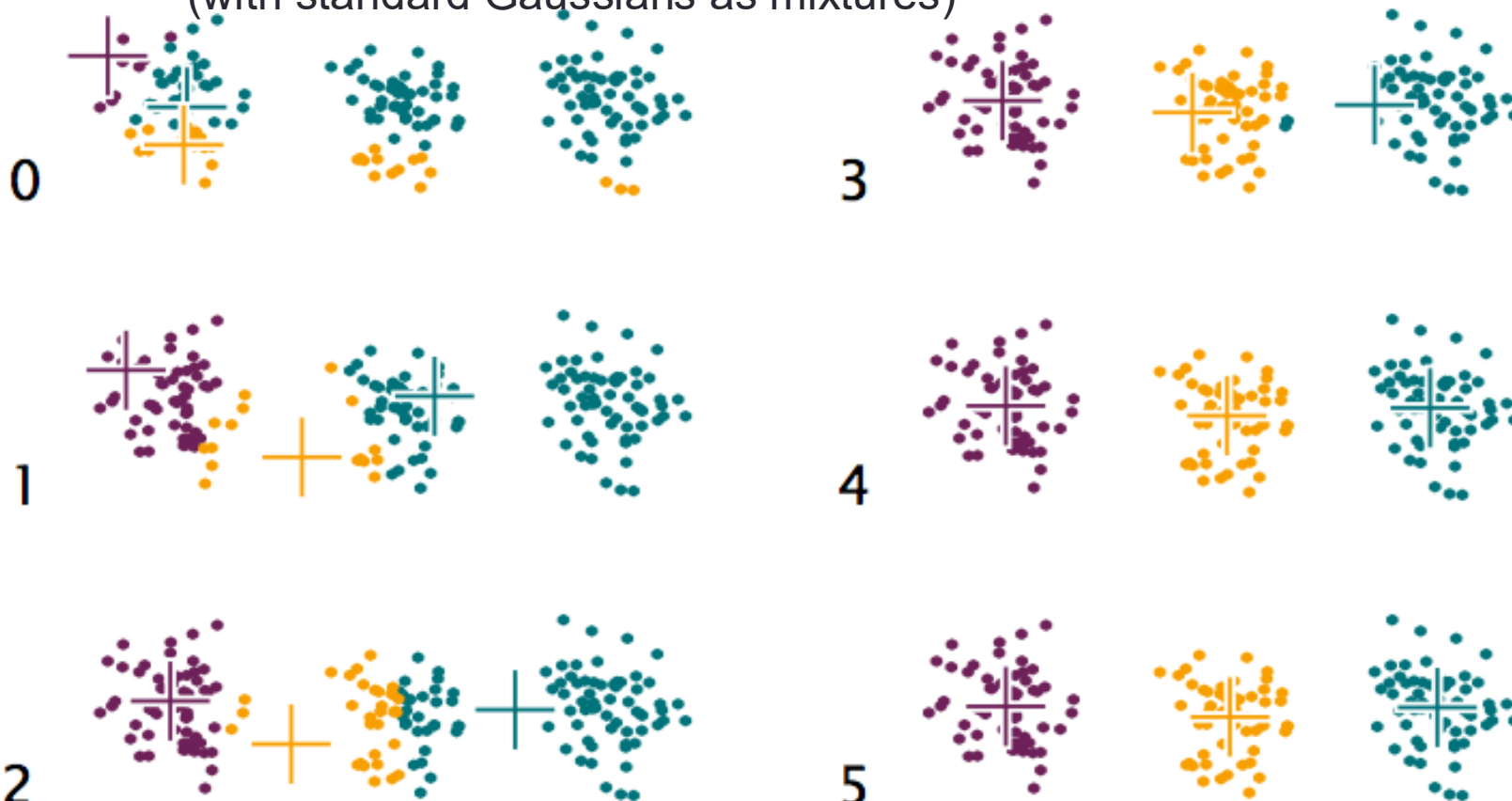
$$\phi_j = \frac{1}{N} \sum_{n=1}^N w_{n,j}$$

$$\mu_j = \frac{\sum_{n=1}^N w_{n,j} x_n}{\sum_{n=1}^N w_{n,j}}$$

$$\sigma_j^2 = \frac{\sum_{n=1}^N w_{n,j} (x_n - \mu_j)^2}{\sum_{n=1}^N w_{n,j}}$$

K-mean vs EM

EM on GMM can be considered as EM with soft labels
(with standard Gaussians as mixtures)



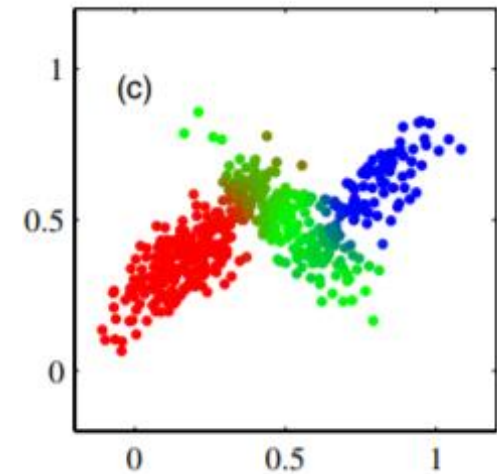
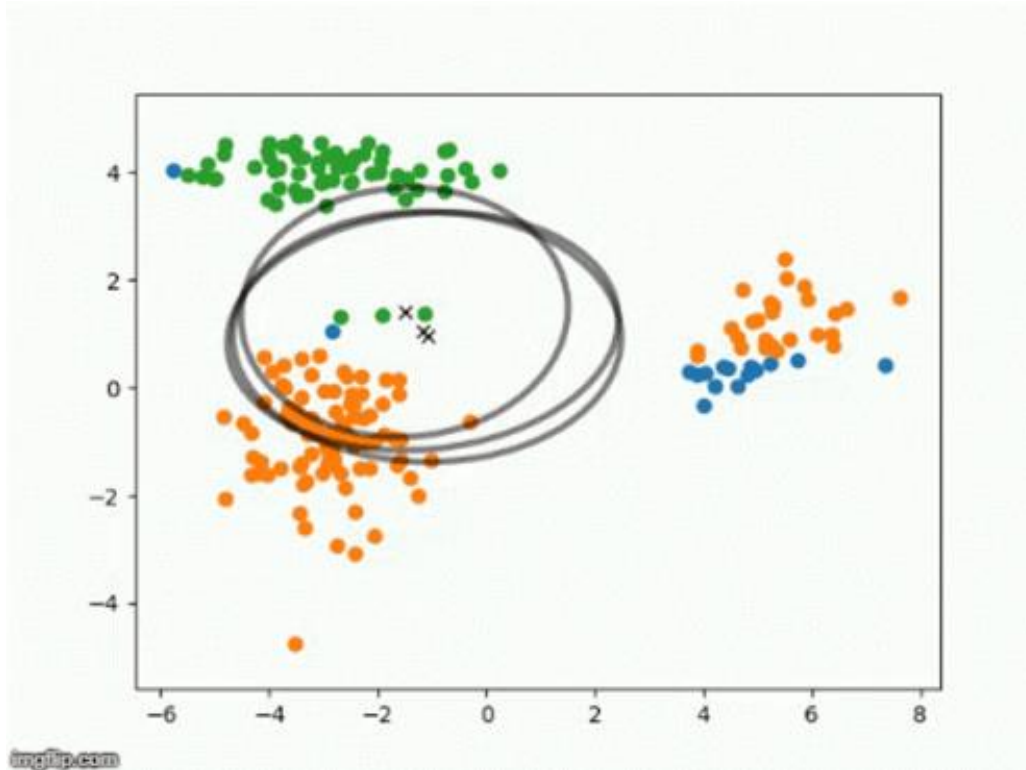
K-mean clustering

- Task: cluster data into groups
- K-mean algorithm
 - **Initialization**: Pick K data points as cluster centers
 - **Assign**: Assign data points to the closest centers
 - **Update**: Re-compute cluster center
 - **Repeat**: Assign and Update

EM algorithm for GMM

- Task: cluster data into Gaussians
- EM algorithm
 - **Initialization**: Randomly initialize parameters Gaussians
 - **Expectation**: Assign data points to the closest Gaussians
 - **Maximization**: Re-compute Gaussians parameters according to assigned data points
 - **Repeat**: Expectation and Maximization
- Note: assigning data points is actually a soft assignment (with probability)

K-mean vs EM

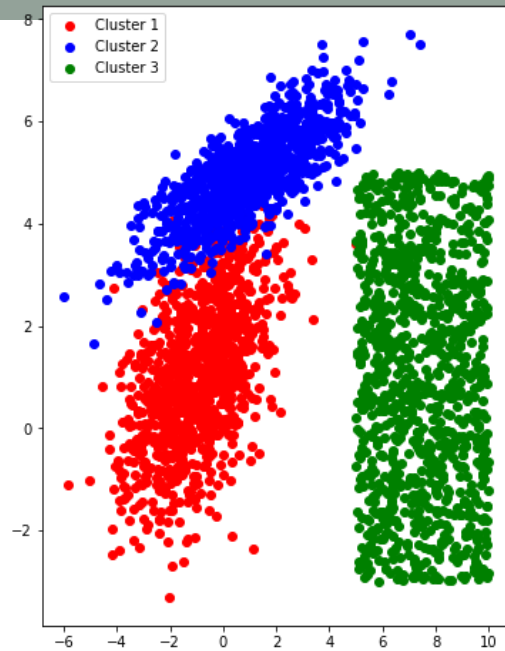


<https://towardsdatascience.com/gaussian-mixture-models-vs-k-means-which-one-to-choose-62f2736025f0>

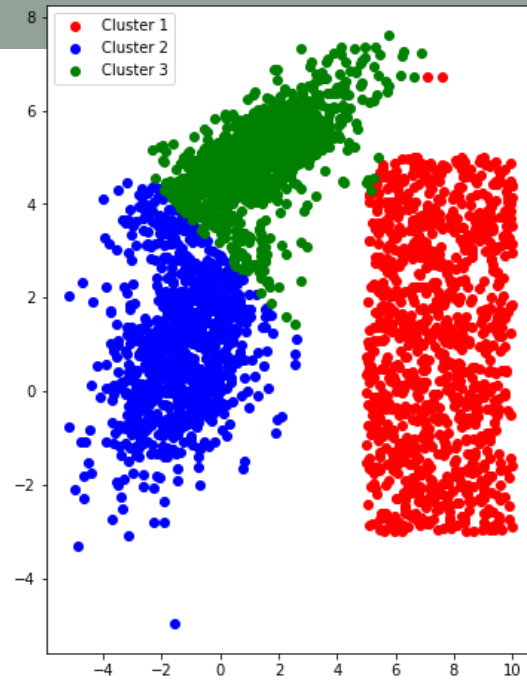
EM/GMM notes

- Converges to local maxima (maximizing likelihood)
 - Just like k-means, need to try different initialization points
- EM always improve the likelihood for each iteration
 - Stops EM when likelihood changes $<$ threshold
- Just like k-means some centroid can get stuck with one sample point and no longer moves
 - For EM on GMM this cause variance to go to 0...
 - Introduce variance floor (minimum variance a Gaussian can have)
- Tricks to avoid bad local maxima
 - Starts with 1 Gaussian
 - Split the Gaussians according to the direction of maximum variance
 - Repeat until arrive at k Gaussians
 - Does not guarantee global maxima but works well in practice

True labels

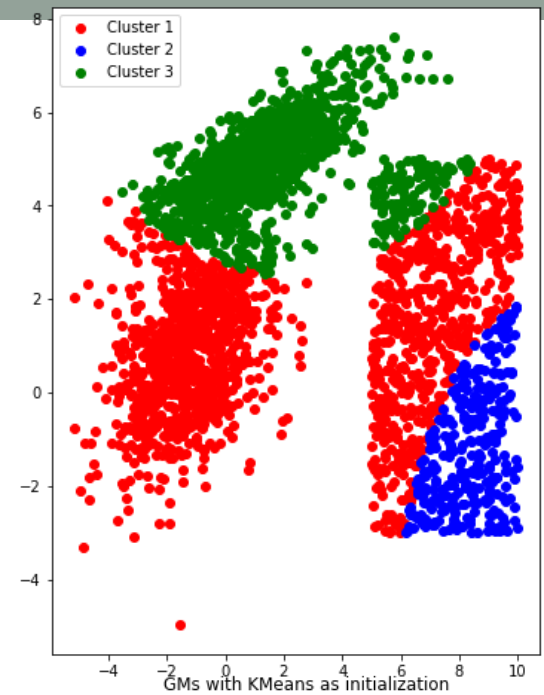


KMeans

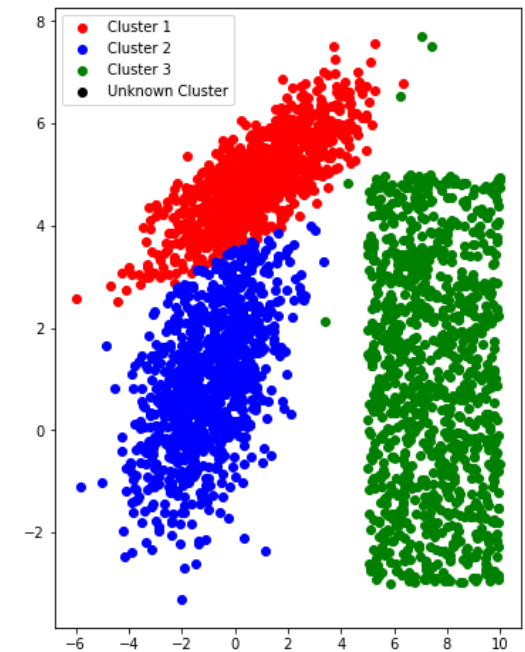


GMs

47

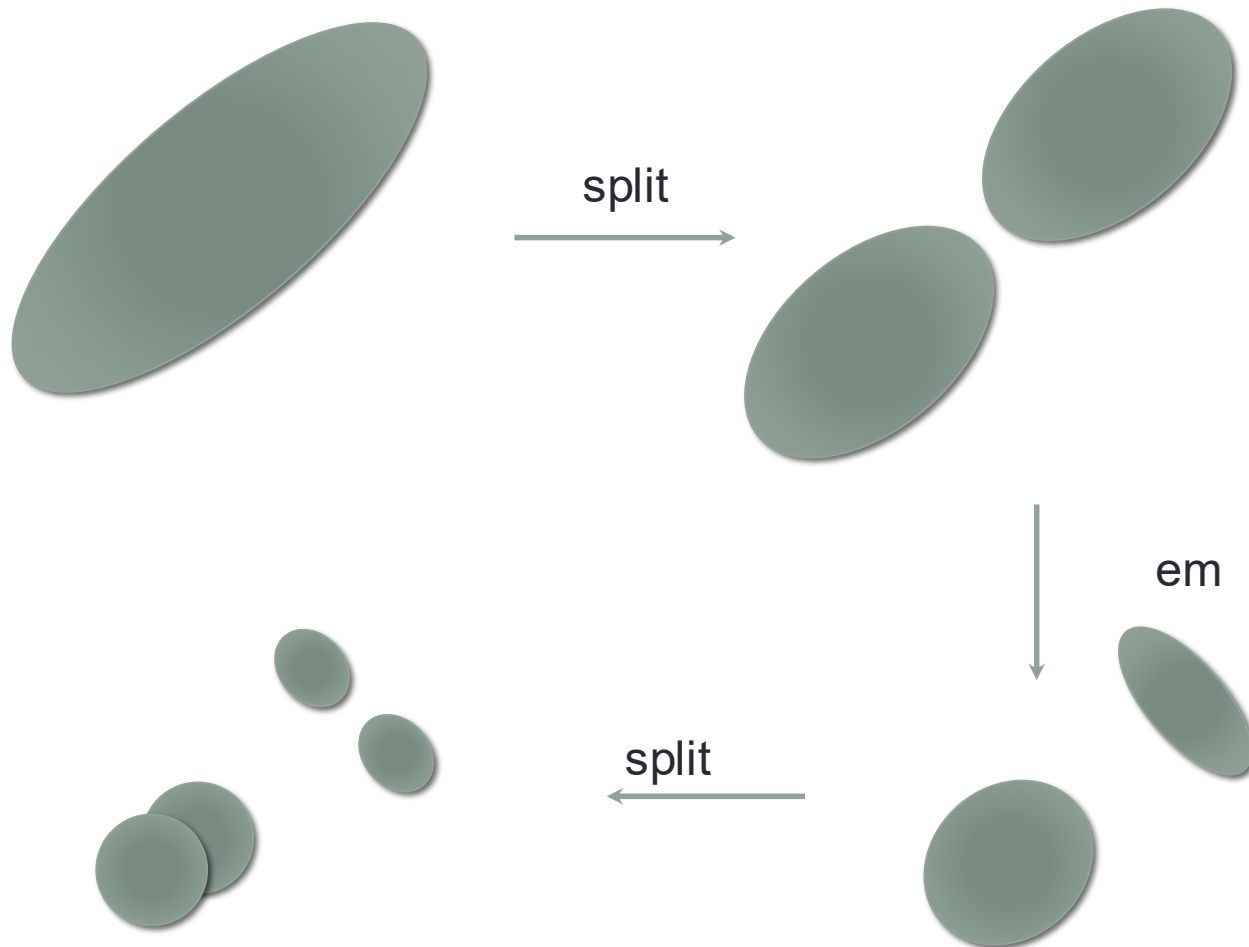


GMs with KMeans as initialization



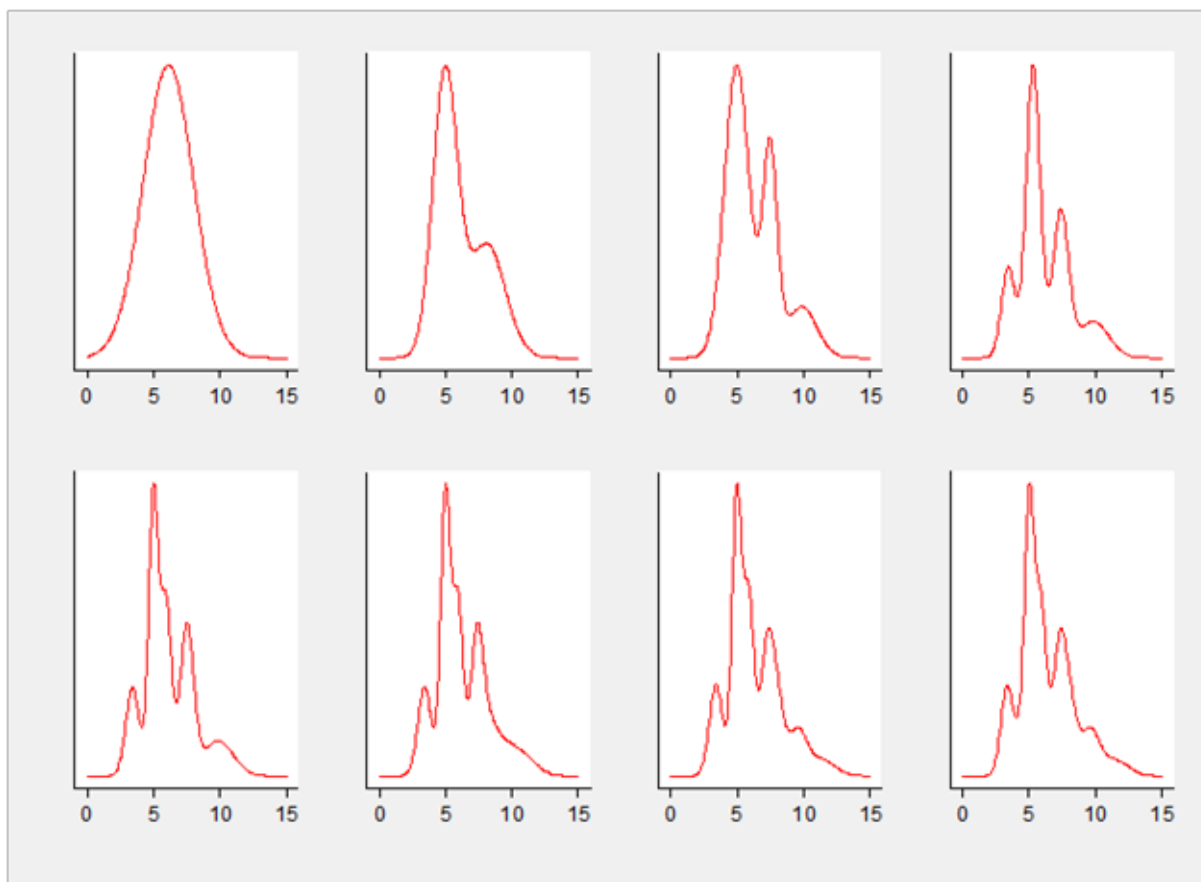
<https://towardsdatascience.com/gaussian-mixture-models-vs-k-means-which-one-to-choose-62f2736025f0>

Gaussian splitting



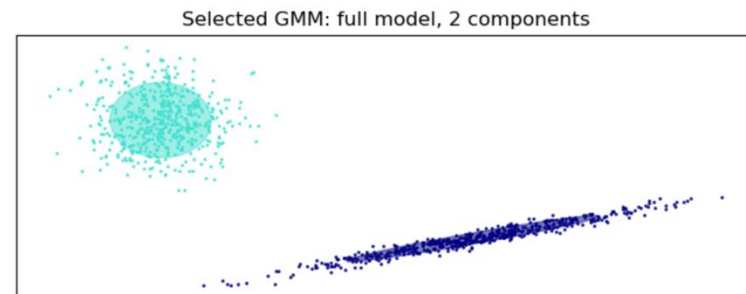
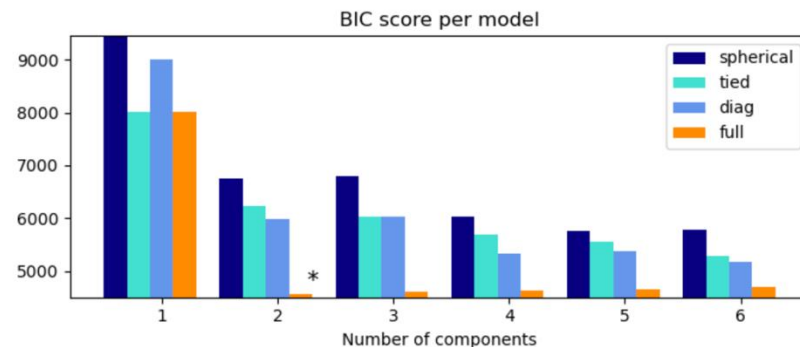
Picking the amount of Gaussians

- As we increase K , the likelihood will keep increasing
- More mixtures \rightarrow more parameters \rightarrow overfits



Picking the amount of Gaussians

- Need a measure of goodness (like Elbow method in k-mean)
- Bayesian Information Criterion (BIC)
- Penalize the log likelihood from the data by the number of parameters in the model
 - $-2 \log L + t \log (n)$
 - t = number of parameters in the model
 - n = number of data points
- We want to minimize BIC



BIC is bad use cross validation!

- Just like how I don't recommend using elbow method for clustering
- BIC is bad use cross validation!
- Test on the goal of your model

Latent variables?

EM is all about problem formulation. You can solve the same task with different formulations.

Latent variable considerations

- Imaginary quantity meant to provide a simplified view of the process
 - GMM mixtures. Speech recognizer states. Customer segmentation.
- Real-world thing, but impossible to directly measure
 - Cause of a disease. Temperature of a star.
- Real-world thing, that is not measured because of noise/faulty sensors

Latent variables?

- **Discrete latent variables**: clusters/partitions data into subgroups
- **Continuous latent variables**: can be used for dimensionality reduction (factor analysis, etc)

EM usage examples

Image segmentation with GMM EM

- D - $\{r, g, b\}$ value at each pixel
- Latent : segment where each pixel comes from
- Hyperparameters: number of mixtures (K), initial values

input

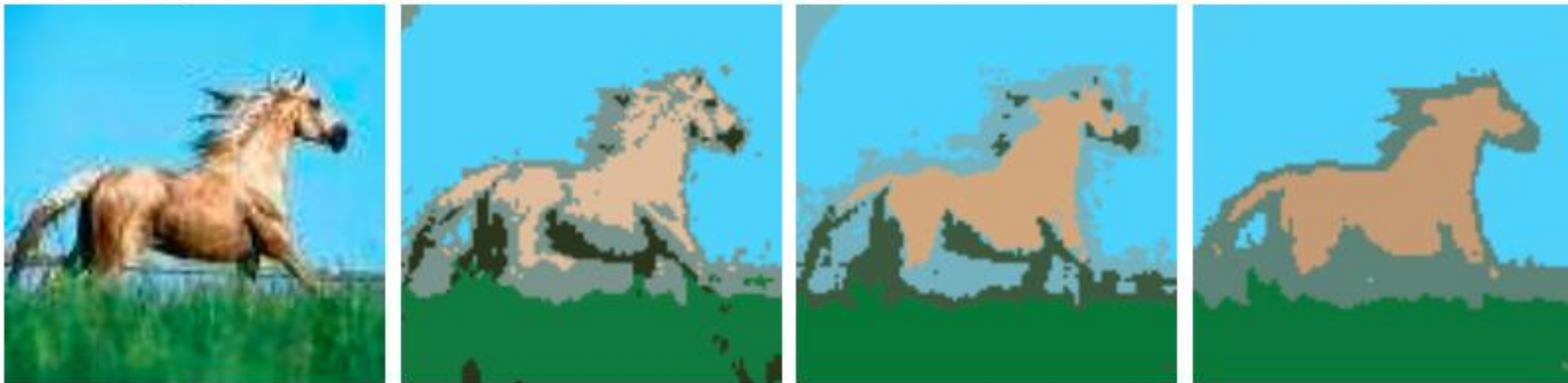


Image segmentation with GMM EM



Fig. 1. Original images: (a) flower, (b) tiger, (c) bear

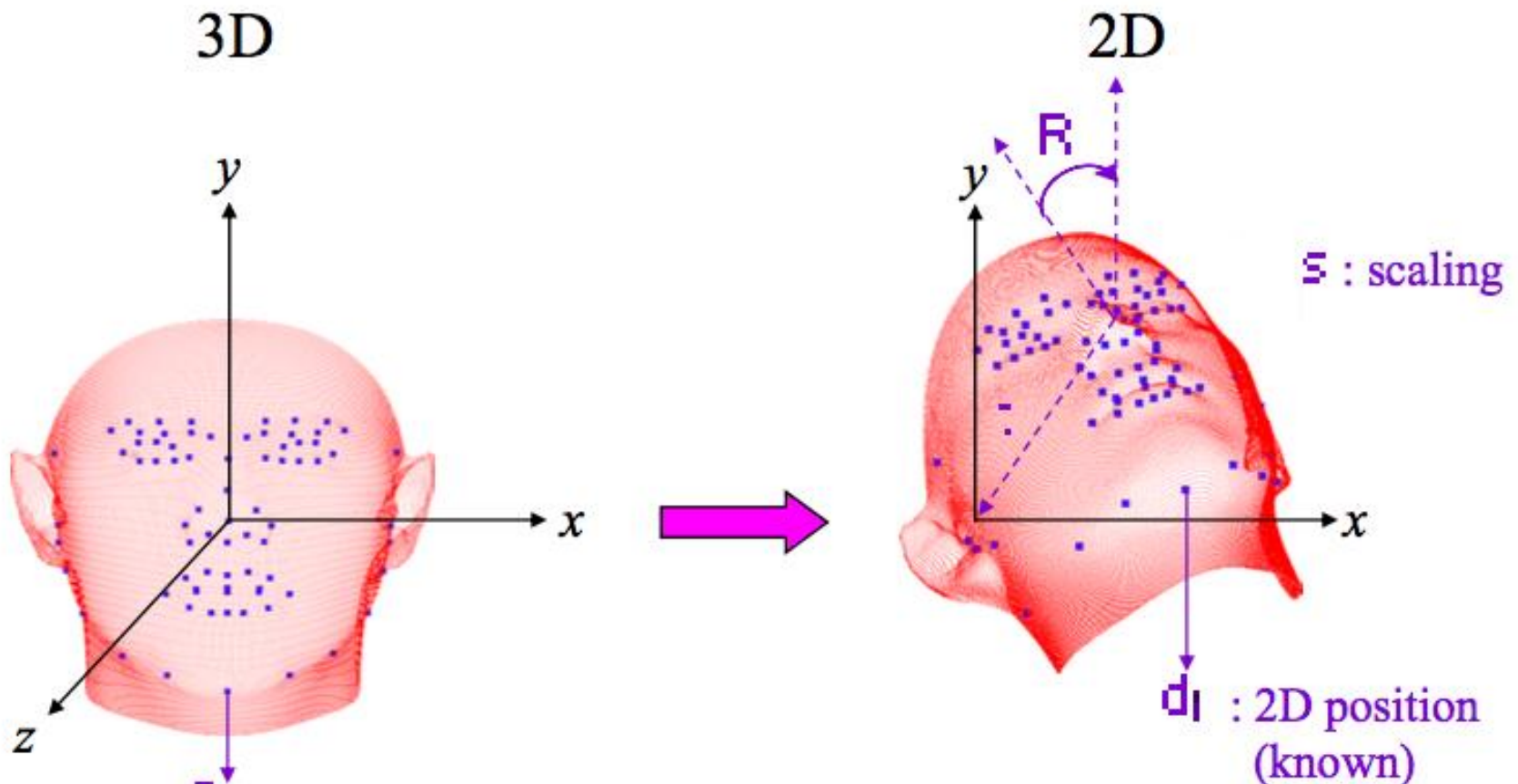


Fig. 2. Segmentation results ($M = 2$)



Fig. 3. Segmentation results ($M = 5$)

Face pose estimation (estimate 3d coordinates from 2d picture)



Language modeling

THE UNITED STATES CONSTITUTION

We the People of the United States, in Order to form a more perfect Union, establish Justice, insure domestic Tranquility, provide for the common defence, promote the general Welfare, and secure the Blessings of Liberty to ourselves and our Posterity, do ordain and establish this Constitution for the United States of America.

Article I

Section 1.

All legislative Powers herein granted shall be vested in a Congress of the United States, which shall consist of a Senate and House of Representatives.

Section 2.

Clause 1: The House of Representatives shall be composed of Members chosen every second Year by the People of the several States, and the Electors in each State shall have the Qualifications requisite for Electors of the most numerous Branch of the State Legislature.

Clause 2: No Person shall be a Representative who shall not have attained to the Age of twenty-five Years, and seven years a Citizen of the United States, and who shall not, when elected, be an Inhabitant of that State in which he shall be chosen.

Clause 3: Representatives and direct Taxes shall be apportioned among the several States which may be included within the Union, according to their respective Numbers, which shall be determined by adding to the whole Number of free Persons, including those bound to Service for a Term of Years, and excluding Indians not taxed, three fifths of all other Persons. The actual Enumeration shall be made within three Years after the first Meeting of the Congress of the United

Latent variable:
Topic
 $P(\text{word}|\text{topic})$

For examples: see Probabilistic latent semantic analysis

MEME

Multiple EM for Motif Elicitation

From Wikipedia, the free encyclopedia

For other uses, see [MEME \(disambiguation\)](#).

Multiple Expectation maximizations for Motif Elicitation (MEME) is a tool for discovering motifs in a group of related [DNA](#) or [protein](#) sequences.^[1]

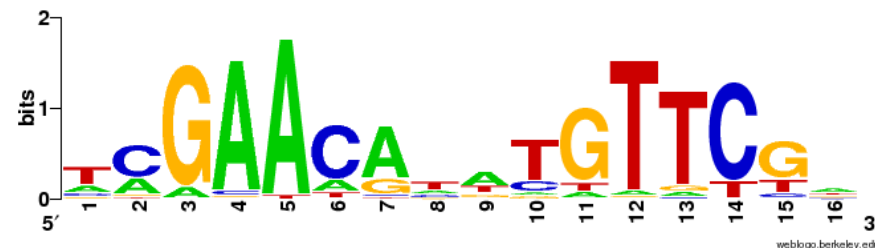
A [motif](#) is a sequence pattern that occurs repeatedly in a group of related protein or DNA sequences and is often associated with some biological function. MEME represents motifs as [position-dependent letter-probability matrices](#) which describe the probability of each possible letter at each position in the pattern. Individual MEME motifs do not contain gaps. Patterns with variable-length gaps are split by MEME into two or more separate motifs.

MEME takes as input a group of DNA or protein sequences (the training set) and outputs as many motifs as requested. It uses statistical modeling techniques to automatically choose the best width, number of occurrences, and description for each motif.

MEME is the first of a collection of tools for analyzing motifs called the [MEME suite](#).

Contents [\[hide\]](#)

- [Definition](#)
- [Use](#)
- [Algorithm components](#)
- [See also](#)
- [References](#)
- [External links](#)



https://en.wikipedia.org/wiki/Multiple_EM_for_Motif_Elicitation

https://en.wikipedia.org/wiki/Position_weight_matrix

Summary

- GMM
 - Mixture of Gaussians
- EM
 - Expectation
 - Maximization

More info and exact proofs

https://seanborman.com/publications/EM_algorithm.pdf

<http://cs229.stanford.edu/summer2019/cs229-notes8.pdf>