

Clustering

Lecture 5: Mixture Model

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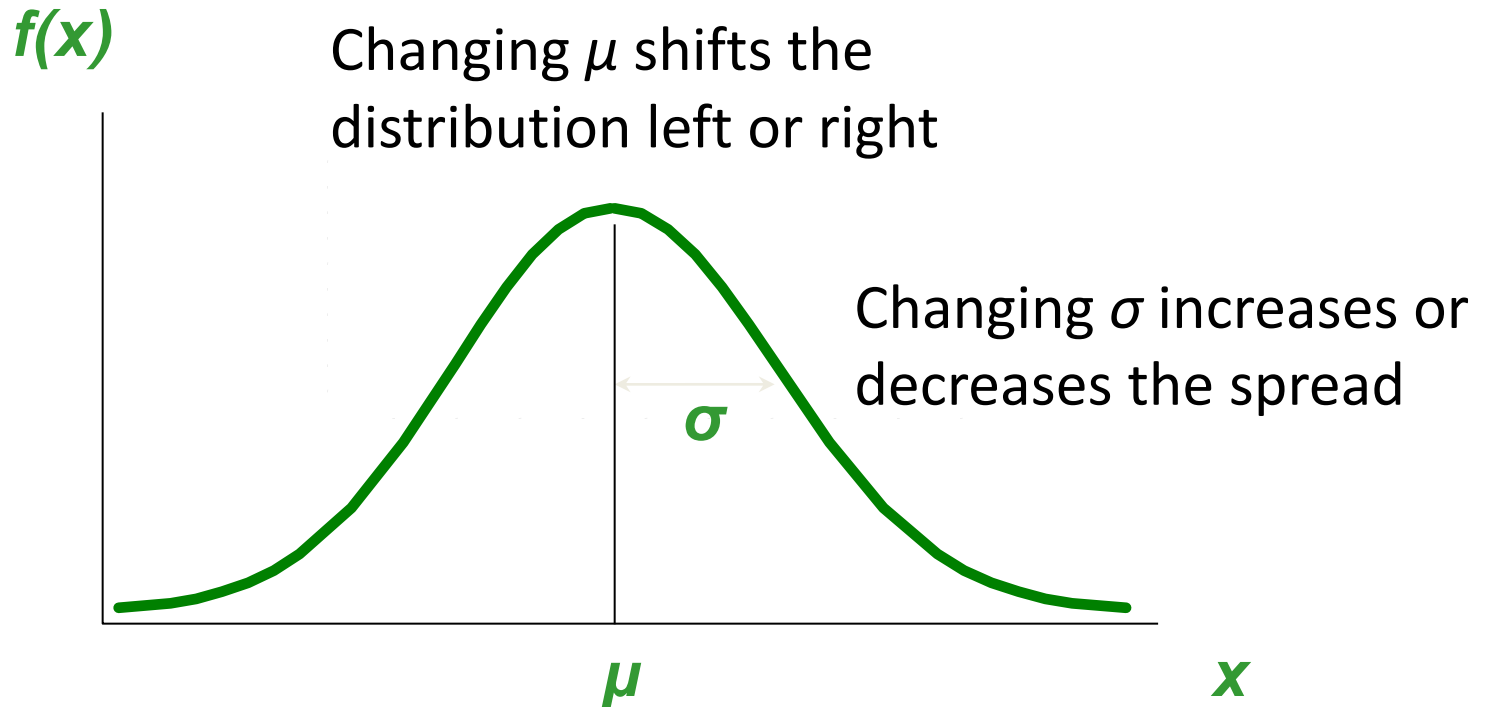
Outline

- **Basics**
 - Motivation, definition, evaluation
- **Methods**
 - Partitional
 - Hierarchical
 - Density-based
 - Mixture model
 - Spectral methods
- **Advanced topics**
 - Clustering ensemble
 - Clustering in MapReduce
 - Semi-supervised clustering, subspace clustering, co-clustering, etc.

Using Probabilistic Models for Clustering

- **Hard vs. soft clustering**
 - Hard clustering: Every point belongs to exactly one cluster
 - Soft clustering: Every point belongs to several clusters with certain degrees
- **Probabilistic clustering**
 - Each cluster is mathematically represented by a parametric distribution
 - The entire data set is modeled by a mixture of these distributions

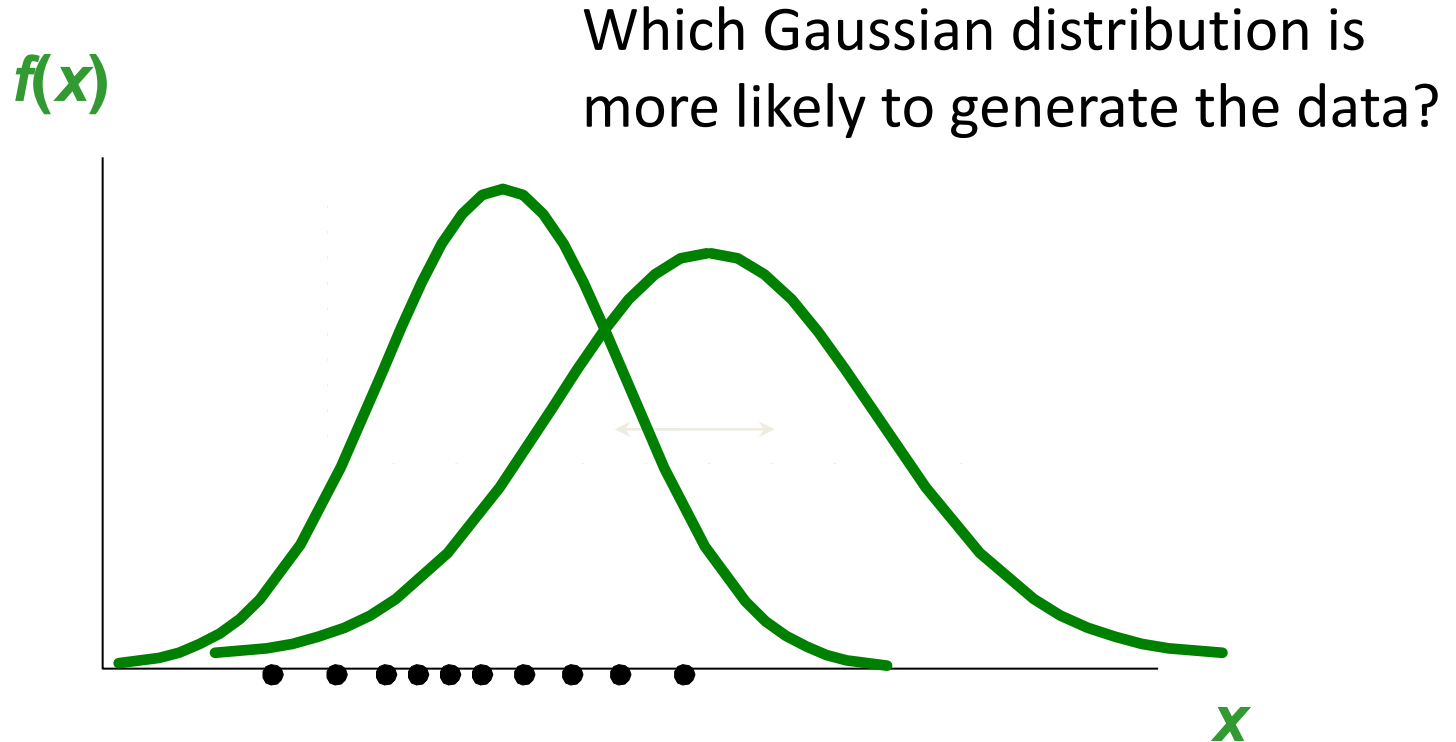
Gaussian Distribution



Probability density function $f(x)$ is a function of x given μ and σ

$$N(x | \mu, \sigma^2) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left(-\frac{1}{2}\left(\frac{x-\mu}{\sigma}\right)^2\right)$$

Likelihood



Define likelihood as a function of μ and σ
given x_1, x_2, \dots, x_n

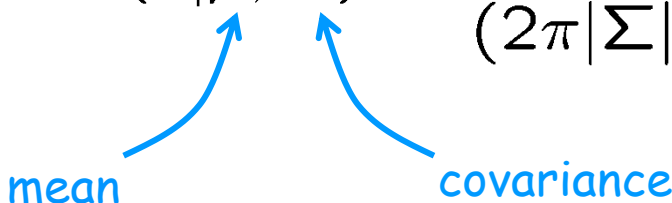
$$\prod_{i=1}^n N(x_i | \mu, \sigma^2)$$

Gaussian Distribution

- Multivariate Gaussian

$$\mathcal{N}(x|\mu, \Sigma) = \frac{1}{(2\pi|\Sigma|)^{1/2}} \exp \left\{ -\frac{1}{2}(x - \mu)^T \Sigma^{-1}(x - \mu) \right\}$$

mean covariance



- Log likelihood

$$L(\mu, \Sigma) = \sum_{i=1}^n \ln N(x_i | \mu, \Sigma) = \sum_{i=1}^n \left(-\frac{1}{2}(x_i - \mu)^T \Sigma^{-1}(x_i - \mu) - \pi \ln |\Sigma| \right)$$

Maximum Likelihood Estimate

- MLE
 - Find model parameters μ, Σ that maximize log likelihood

$$L(\mu, \Sigma)$$

- MLE for Gaussian

$$\hat{\mu} = \frac{1}{n} \sum_{i=1}^n x_i$$

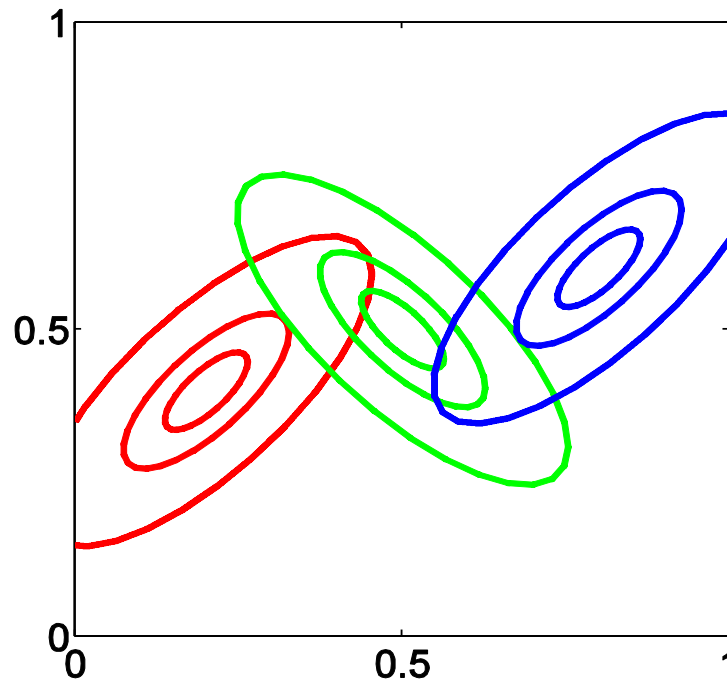
$$\hat{\Sigma} = \frac{1}{n} \sum_{i=1}^n (x_i - \hat{\mu})(x_i - \hat{\mu})^T$$

Gaussian Mixture

- Linear combination of Gaussians

$$p(x) = \sum_{k=1}^K \pi_k \mathcal{N}(x | \mu_k, \Sigma_k) \quad \text{where} \quad \sum_{k=1}^K \pi_k = 1, \quad 0 \leq \pi_k \leq 1$$

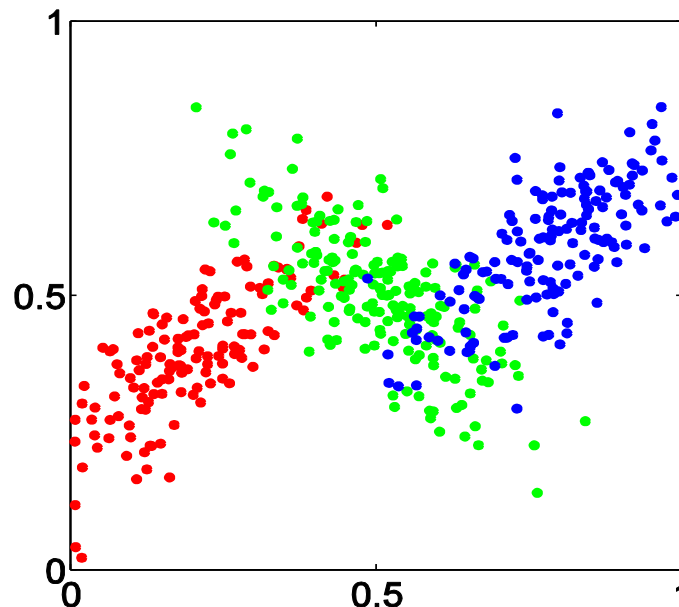
parameters to be estimated



Gaussian Mixture

- To generate a data point:
 - first pick one of the components with probability π_k
 - then draw a sample \mathbf{x}_i from that component distribution
- Each data point is generated by one of K components, a **latent** variable $\mathbf{z}_i = (z_{i1}, \dots, z_{iK})$ is associated with each \mathbf{x}_i

$$\sum_{k=1}^K z_{ik} = 1 \text{ and } p(z_{ik} = 1) = \pi_k$$



Gaussian Mixture

- Maximize log likelihood

$$\ln p(x|\pi, \mu, \Sigma) = \sum_{i=1}^n \ln \left\{ \sum_{k=1}^K \pi_k \mathcal{N}(x_i | \mu_k, \Sigma_k) \right\}$$

- Without knowing values of latent variables, we have to maximize the **incomplete** log likelihood

Expectation-Maximization (EM) Algorithm

- E-step: for given parameter values we can compute the expected values of the latent variables (**responsibilities** of data points)

$$\begin{aligned} r_{ik} \equiv E(z_{ik}) &= p(z_{ik} = 1 | x_i, \pi, \mu, \Sigma) \\ &= \frac{p(z_{ik} = 1) p(x_i | z_{ik} = 1, \pi, \mu, \Sigma)}{\sum_{k=1}^K p(z_{ik} = 1) p(x_i | z_{ik} = 1, \pi, \mu, \Sigma)} \\ &= \frac{\pi_k \mathcal{N}(x_i | u_k, \Sigma_k)}{\sum_{k=1}^K \pi_k \mathcal{N}(x_i | u_k, \Sigma_k)} \end{aligned}$$

- Note that $r_{ik} \in [0, 1]$ instead of $\{0, 1\}$ but we still have $\sum_{k=1}^K r_{ik} = 1$ for all i

Expectation-Maximization (EM) Algorithm

- M-step: maximize the **expected complete** log likelihood

$$E[\ln p(x, z|\pi, \mu, \Sigma)] = \sum_{i=1}^n \sum_{k=1}^K r_{ik} \{\ln \pi_k + \ln \mathcal{N}(x_i|\mu_k, \Sigma_k)\}$$

- Parameter update:

$$\pi_k = \frac{\sum_i r_{ik}}{n} \quad \mu_k = \frac{\sum_i r_{ik} x_i}{\sum_i r_{ik}}$$

$$\Sigma_k = \frac{\sum_i r_{ik} (x_i - \mu_k)(x_i - \mu_k)^T}{\sum_i r_{ik}}$$

EM Algorithm

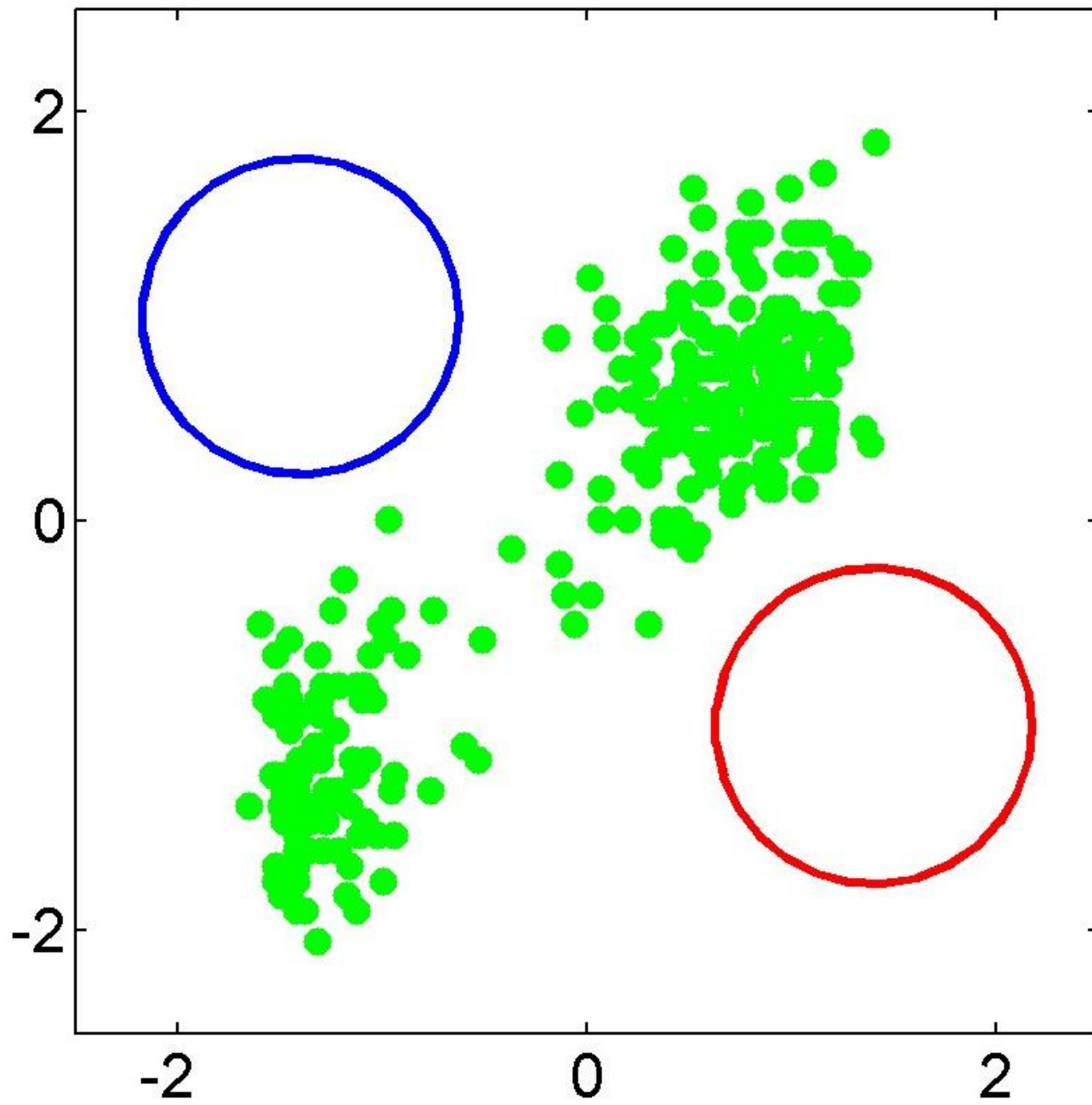
- Iterate E-step and M-step until the log likelihood of data does not increase any more.
 - Converge to **local optimal**
 - Need to restart algorithm with different initial guess of parameters (as in *K*-means)

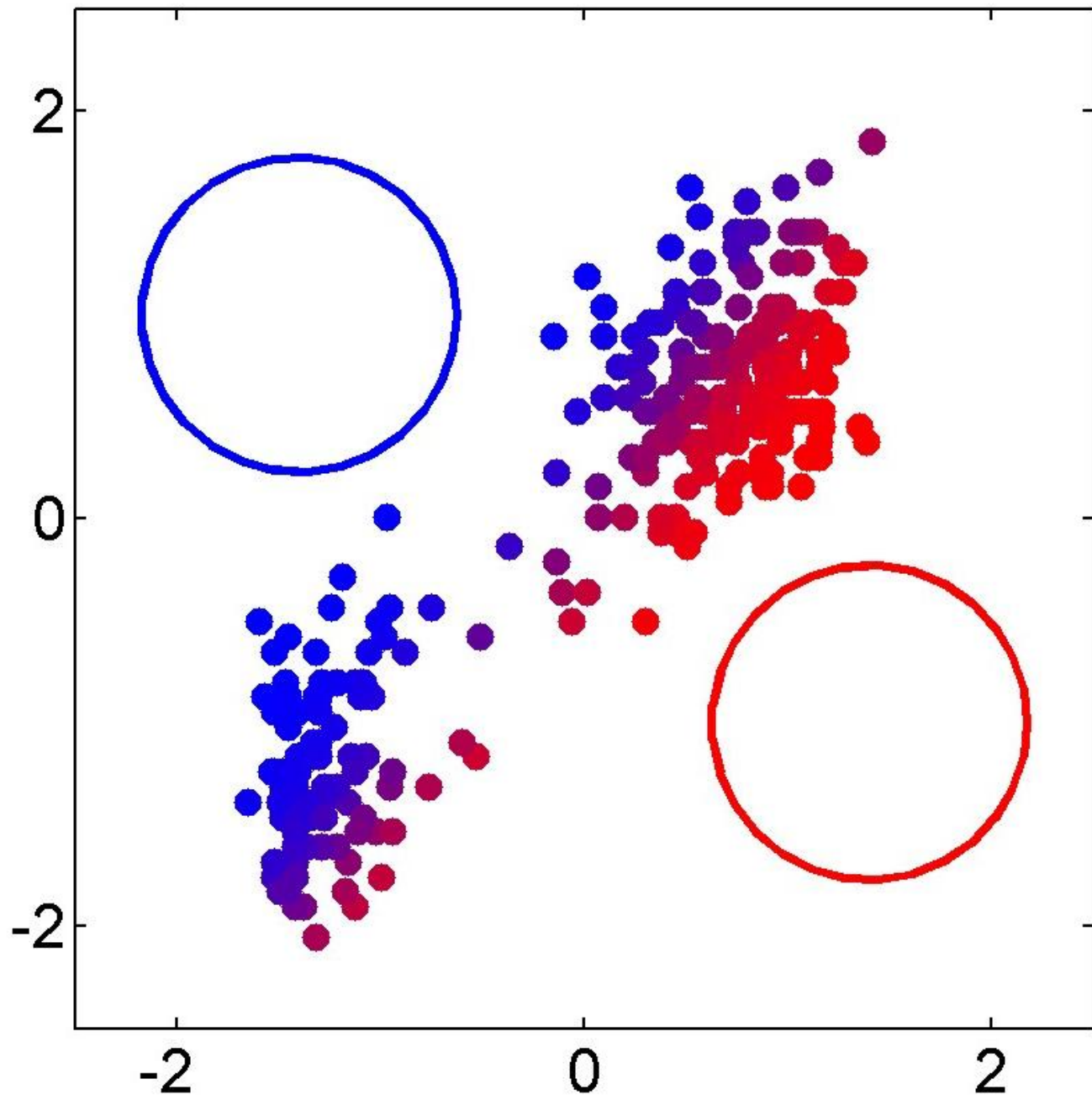
- Relation to *K*-means

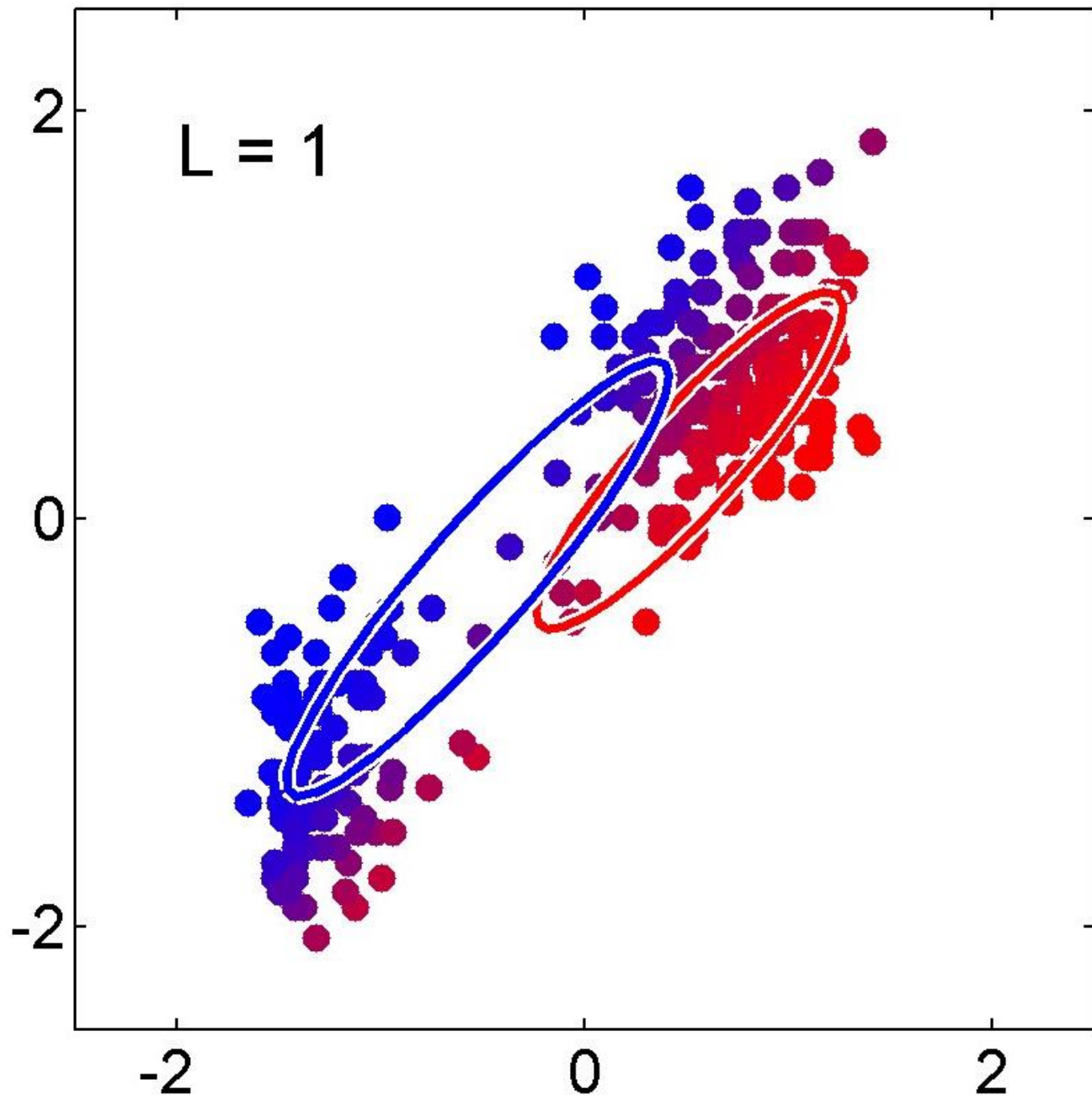
- Consider GMM with common covariance

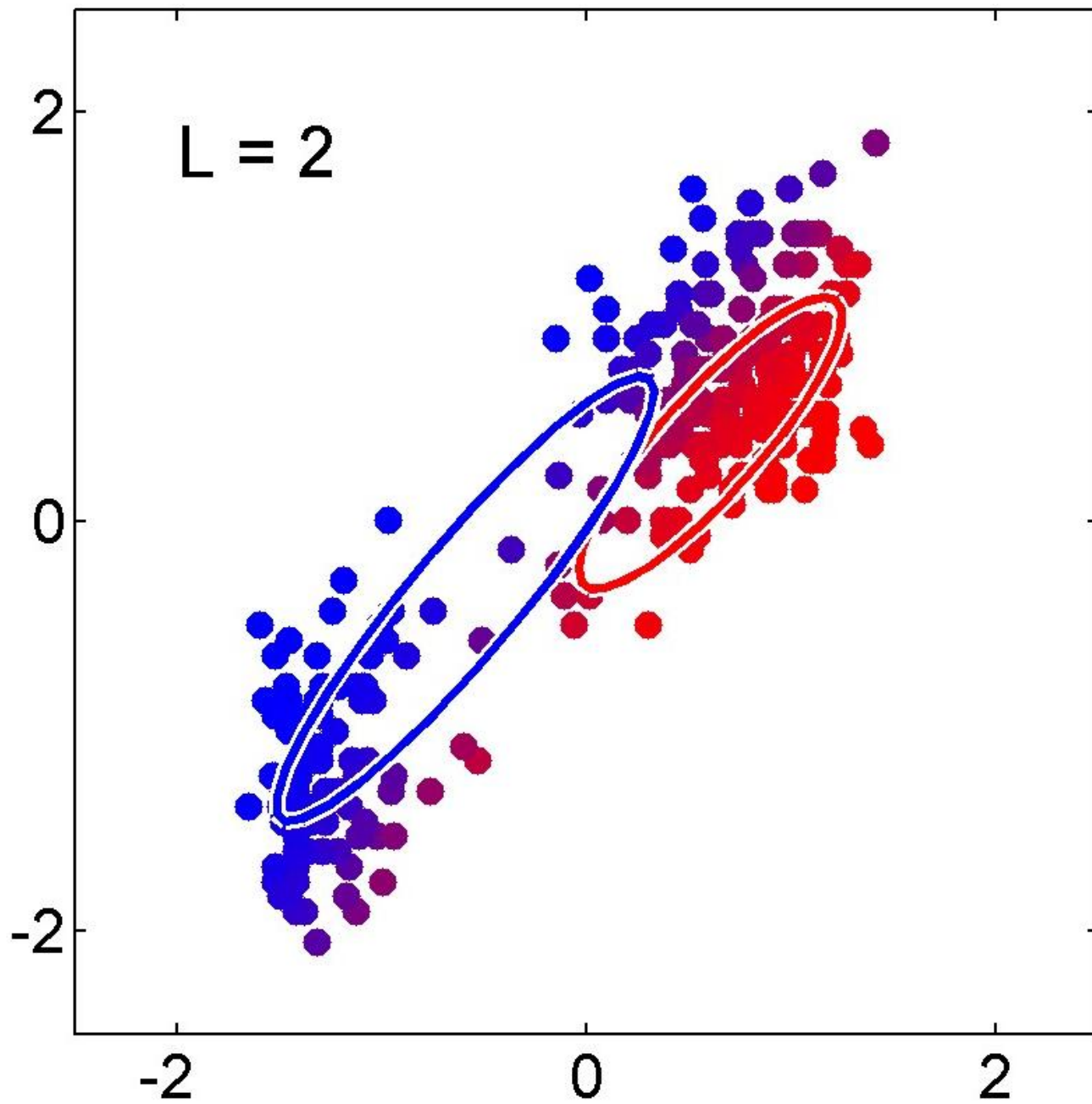
$$\Sigma_k = \delta^2 I$$

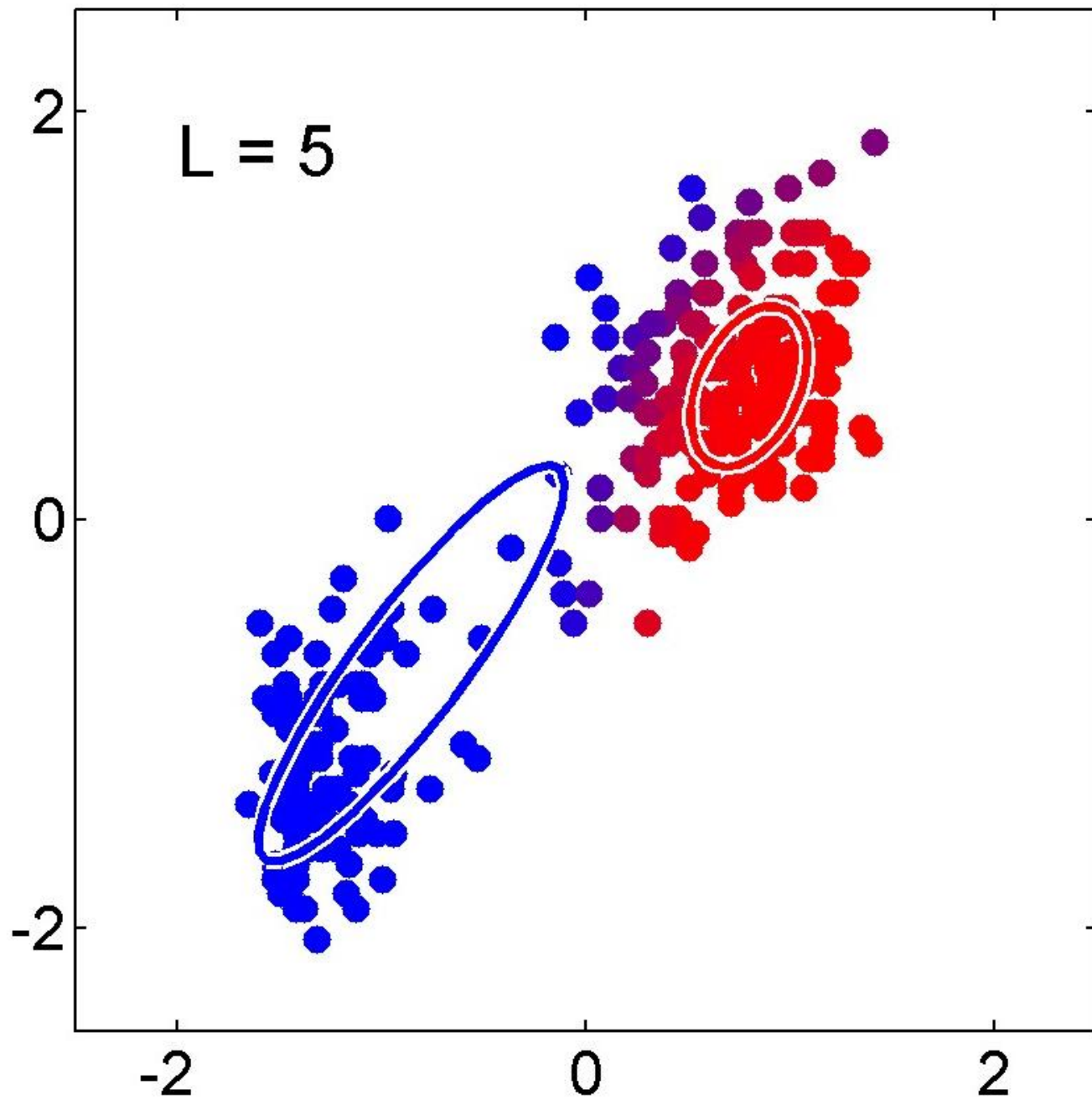
- As $\delta^2 \rightarrow 0$, $r_{ik} \rightarrow 0$ or 1 , two methods coincide

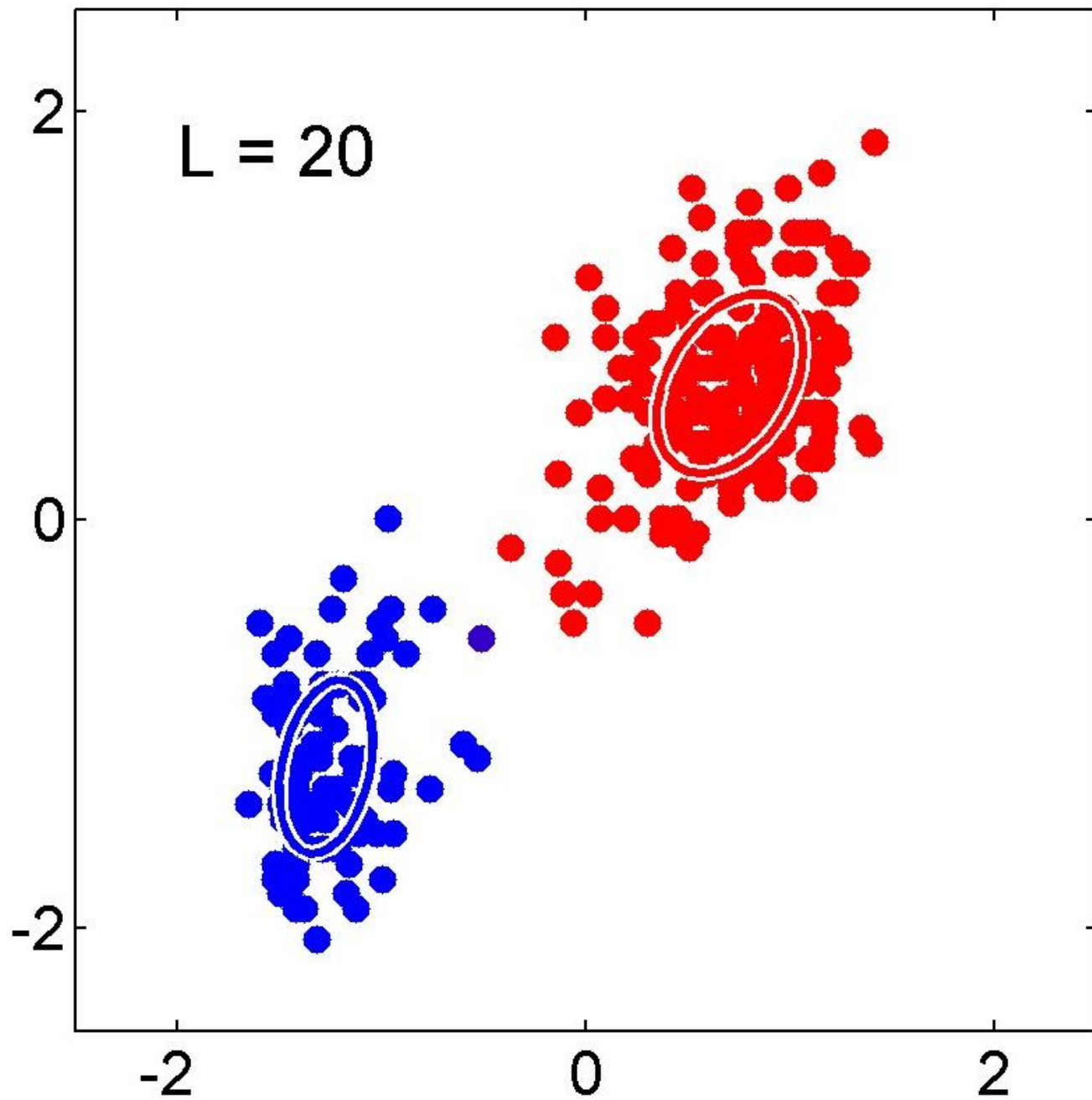












K-means vs GMM

- Objective function
 - Minimize sum of squared Euclidean distance
 - Can be optimized by an EM algorithm
 - E-step: assign points to clusters
 - M-step: optimize clusters
 - Performs hard assignment during E-step
 - Assumes spherical clusters with equal probability of a cluster
- Objective function
 - Maximize log-likelihood
 - EM algorithm
 - E-step: Compute posterior probability of membership
 - M-step: Optimize parameters
 - Perform soft assignment during E-step
 - Can be used for non-spherical clusters
 - Can generate clusters with different probabilities

Mixture Model

- **Strengths**

- Give probabilistic cluster assignments
- Have probabilistic interpretation
- Can handle clusters with varying sizes, variance etc.

- **Weakness**

- Initialization matters
- Choose appropriate distributions
- Overfitting issues

Take-away Message

- Probabilistic clustering
- Maximum likelihood estimate
- Gaussian mixture model for clustering
- EM algorithm that assigns points to clusters and estimates model parameters alternatively
- Strengths and weakness