

COMS 4771
Assignment Project Exam Help

Clustering
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Nakul Verma

Supervised Learning

Data: $(\vec{x}_1, y_1), (\vec{x}_2, y_2), \dots \in \mathcal{X} \times \mathcal{Y}$

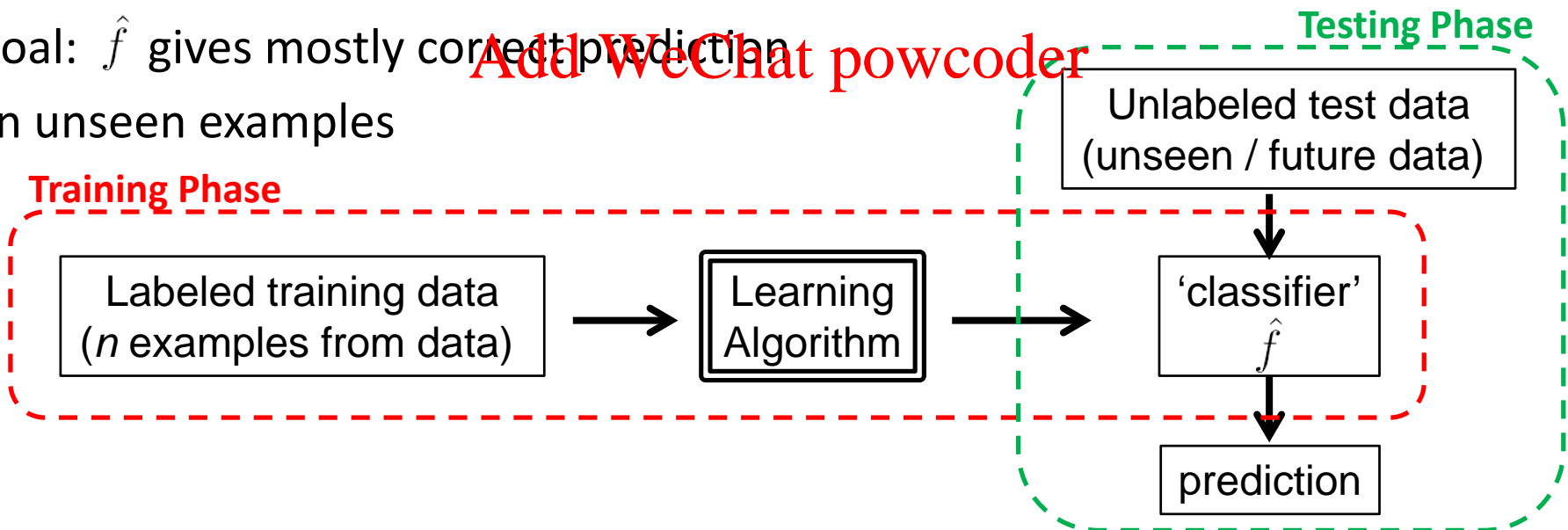
Supervised learning

Assumption: there is a (relatively simple) function $f^* : \mathcal{X} \rightarrow \mathcal{Y}$

such that $f^*(\vec{x}_i) = y_i$ for most i

Learning task: given n examples from the data, find an approximation $\hat{f} \approx f^*$

Goal: \hat{f} gives mostly correct prediction on unseen examples



Unsupervised Learning

Data: $\vec{x}_1, \vec{x}_2, \dots \in \mathcal{X}$

Unsupervised learning

Assumption: there is an underlying structure in \mathcal{X}

Learning task: discover the structure given n examples from the data

Goal: come up with the summary of the data using the discovered structure

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Partition the data into meaningful structures

clustering

Find a low-dimensional representation that retains important information, and suppresses irrelevant/noise information

Dimensionality
reduction

Let's take a closer look using an example...

Example: Handwritten digits revisited

Handwritten digit data, but with no labels

0 1 2 3 4 5 6 7 8 9
0 1 2 3 4 5 6 7 8 9
0 1 2 3 4 5 6 7 8 9
0 1 2 3 4 5 6 7 8 9
0 1 2 3 4 5 6 7 8 9

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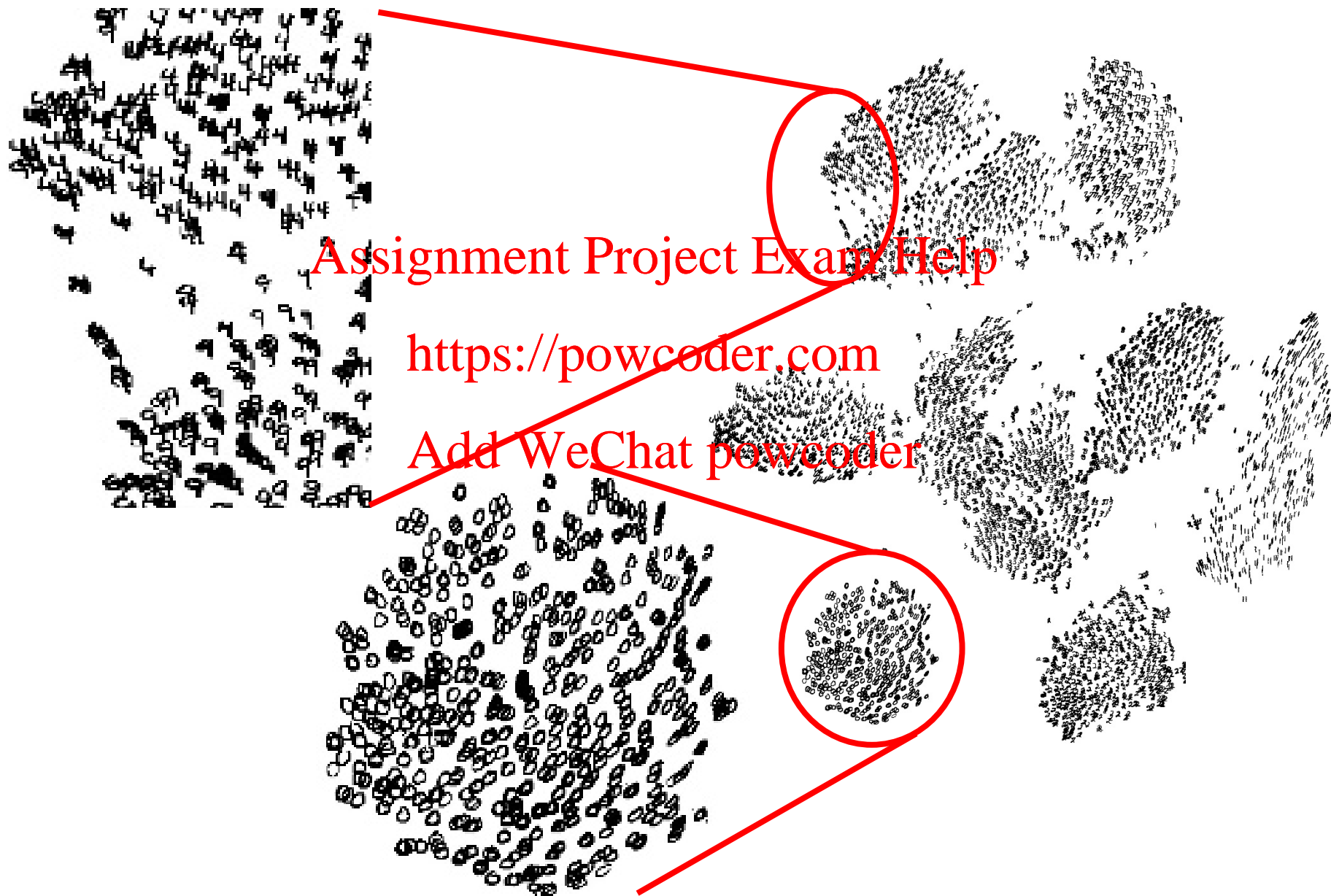
What can we do?

- Suppose know that there are 10 groupings, can we *find the groups*?
- What if we don't know there are 10 groups?
- How can we *discover/explore* other structure in such data?

A 2D visualization of digits dataset



Handwritten digits visualization



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Grouping The Data, aka Clustering

Data: $\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \in \mathcal{X}$

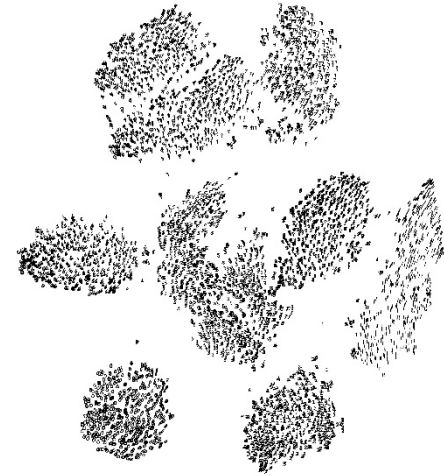
Given: known target number of groups k

Output: Partition $\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n$ into k groups

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**This is called the clustering problem,
also known as unsupervised classification, or quantization**

k-means

Given: data $\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \in \mathbf{R}^d$, and intended number of groupings k

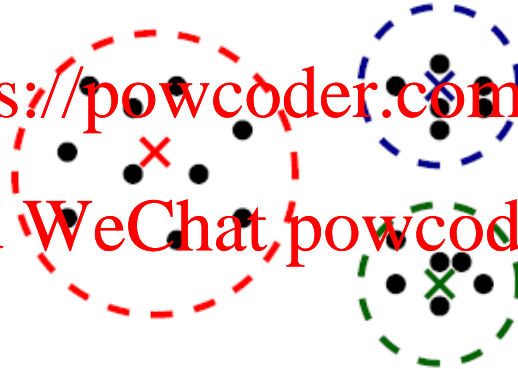
Idea:

find a set of representatives $\vec{c}_1, \vec{c}_2, \dots, \vec{c}_k$ such that data is **close to** some representative

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Optimization:

$$\text{minimize}_{c_1, \dots, c_k} \left[\sum_{i=1}^n \min_{j=1, \dots, k} \|\vec{x}_i - \vec{c}_j\|^2 \right]$$

*Unfortunately this is NP-hard
Even for $d=2$ and $k=2$*

How do we optimize this?

*How do we solve for
 $d=1$ or $k=1$ case?*

Algorithm to approximate k -means

Given: data $\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \in \mathbf{R}^d$, and intended number of groupings k

Alternating optimization algorithm:

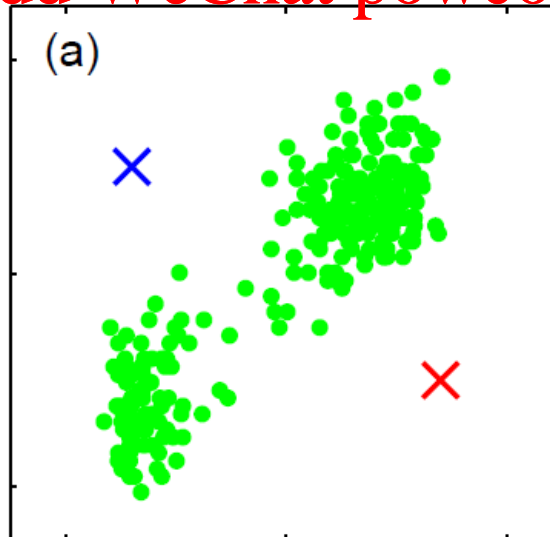
- Initialize cluster centers $\vec{c}_1, \vec{c}_2, \dots, \vec{c}_k$ (say randomly)
- Repeat till no more changes occur
 - Assign data to its closest center (this creates a partition) (assume centers are fixed)
 - Find the optimal centers $\vec{c}_1, \vec{c}_2, \dots, \vec{c}_k$ (assuming the data partition is fixed)

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Demo:



Algorithm to approximate k -means

Given: data $\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \in \mathbf{R}^d$, and intended number of groupings k

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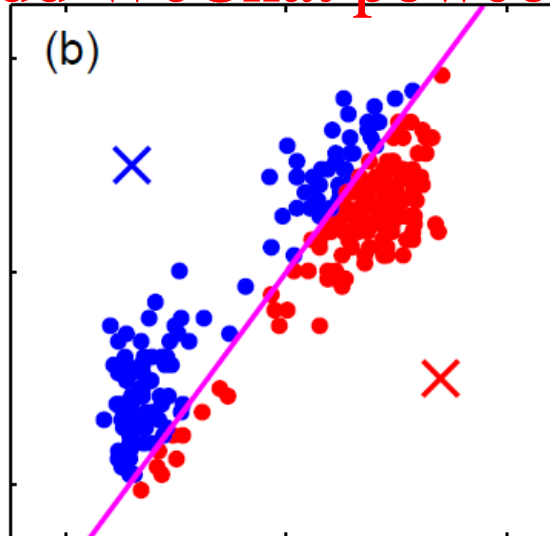
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Demo:



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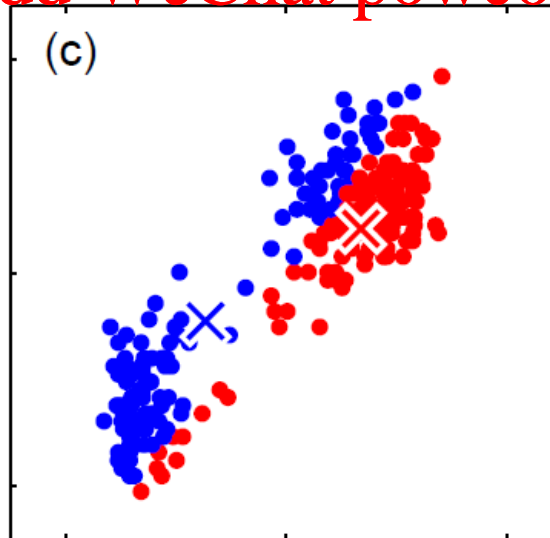
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Demo:



Algorithm to approximate k -means

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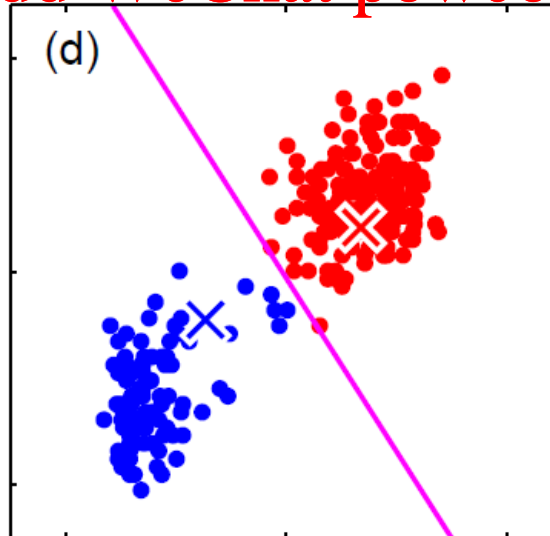
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Demo:



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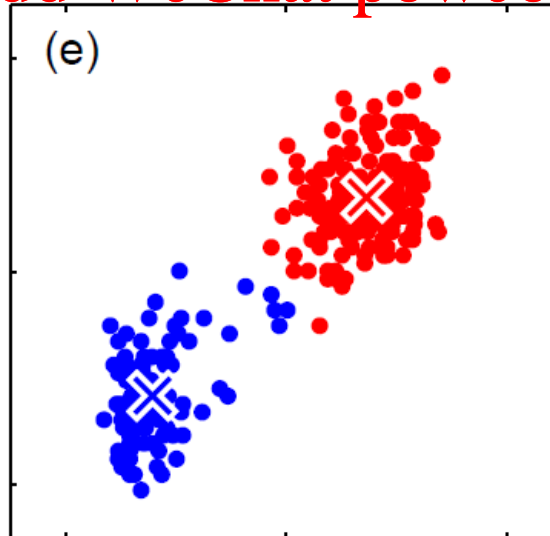
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Demo:



Algorithm to approximate k -means

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Alternating optimization algorithm:

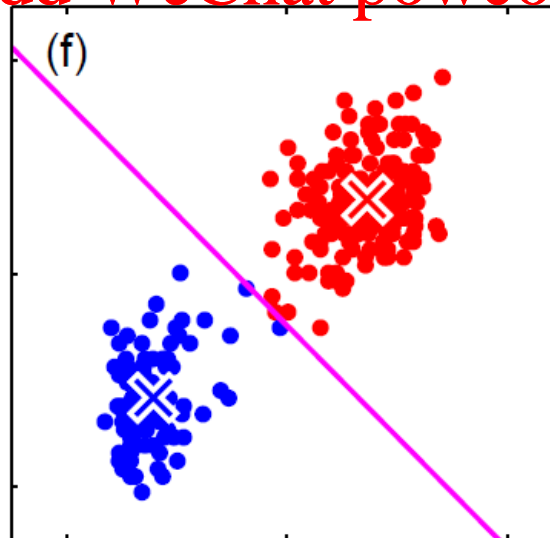
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Demo:



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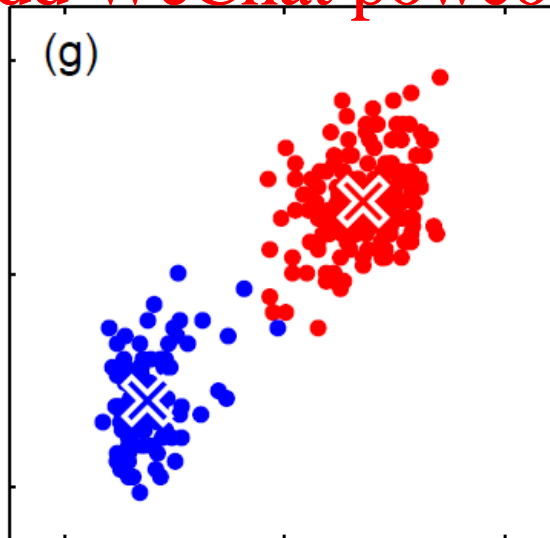
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Demo:



k -means

Some properties of this alternating updates algorithm:

- The approximation can be arbitrarily bad, compared to the best cluster assignment!

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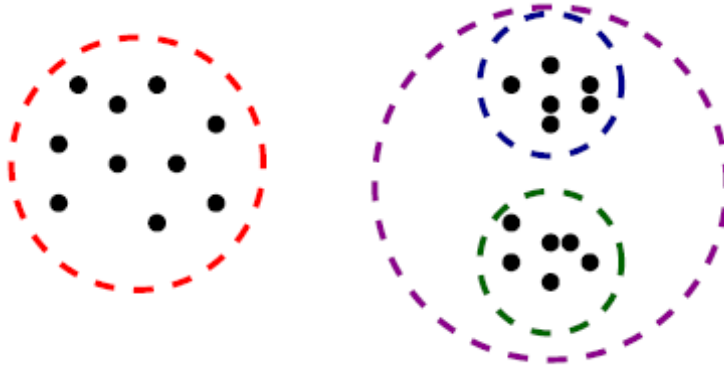
- Performance quality heavily dependent on the initialization!

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k -means:

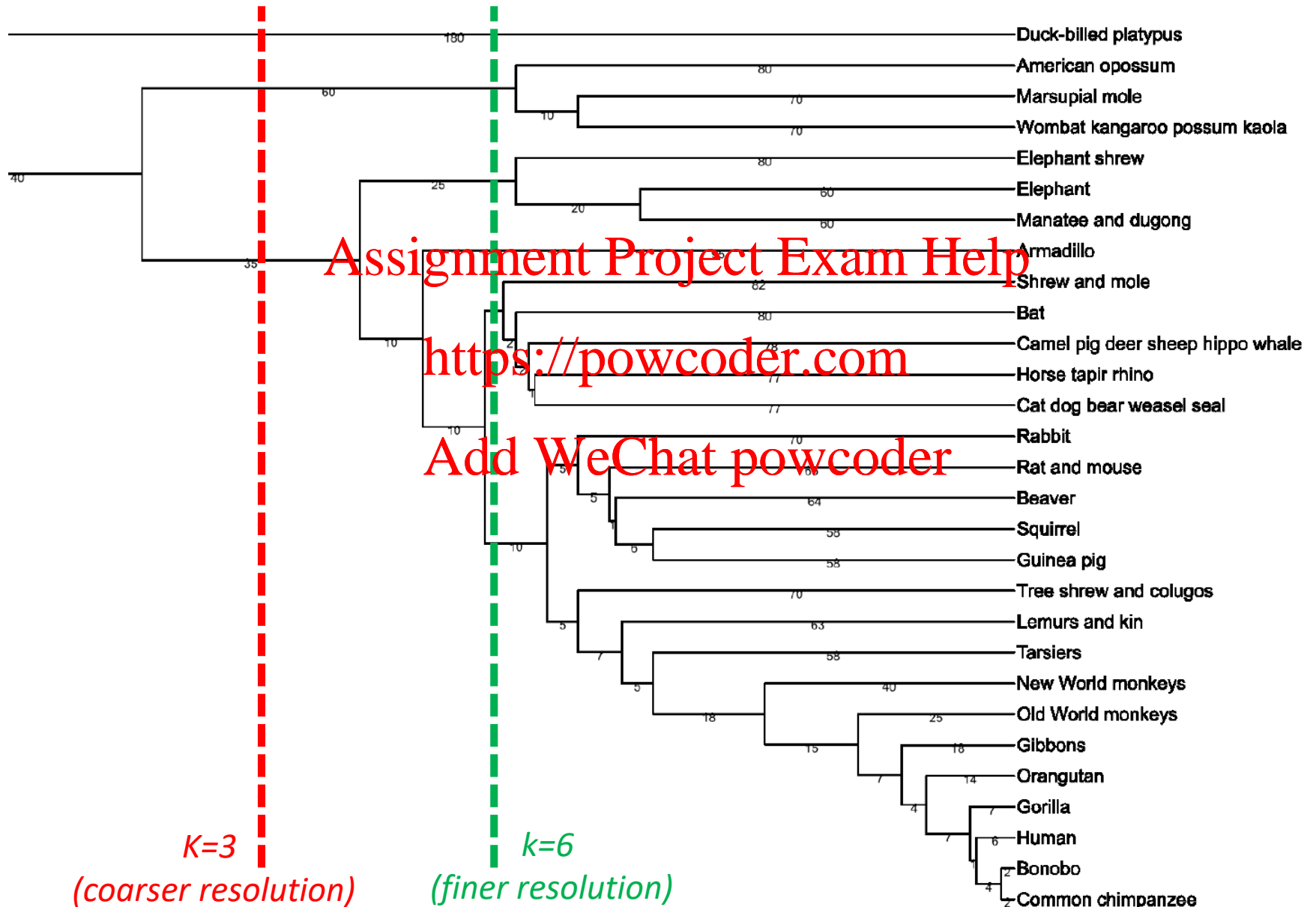
- How to select k ?



is the right $k=2$ or $k=3$?

Solution: encode clustering
for all values of k !
(hierarchical clustering)

Example: Clustering Without Committing to k



Hierarchical Clustering

Two approaches:

Top Down (divisive):

- Partition data into two groups (say, by k-means, with $k=2$)
- Recurse on each part
- Stop when cannot partition data anymore (ie single points left)

Bottom Up (agglomerative):

- Start by each data sample as its own cluster (so initial number of clusters is n)
- Repeatedly merge “closest” pair of clusters
- Stop when only one cluster is left

Clustering via Probabilistic Mixture Modeling

Alternative way to cluster data:

Given: $\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \in \mathbf{R}^d$ and number of intended number of clusters k .

Assume a joint probability distribution (X, C) over the joint space $\mathbf{R}^d \times [k]$

$$C \sim \begin{pmatrix} \pi_1 \\ \vdots \\ \pi_k \end{pmatrix} \quad \text{Discrete distribution over the clusters } P[C=i] = \pi_i$$

$$X|C = i \sim \text{Some multivariate distribution, e.g. } N(\vec{\mu}_i, \Sigma_i)$$

Parameters: $\theta = (\pi_1, \vec{\mu}_1, \Sigma_1, \dots, \pi_k, \vec{\mu}_k, \Sigma_k)$ *looks familiar?*

Modeling assumption data $(x_1, c_1), \dots, (x_n, c_n)$ i.i.d. from $\mathbf{R}^d \times [k]$

BUT only get to see partial information: x_1, x_2, \dots, x_n (c_1, \dots, c_n hidden!)

Gaussian Mixture Modeling (GMM)

Given: $\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \in \mathbf{R}^d$ and k .

Assume a joint probability distribution (X, C) over the joint space $\mathbf{R}^d \times [k]$

$$C \sim \begin{bmatrix} \pi_1 \\ \vdots \\ \pi_k \end{bmatrix} \quad X|C=i \sim N(\vec{\mu}_i, \Sigma_i) \quad \text{Gaussian Mixture Model}$$

$\theta = (\pi_1, \vec{\mu}_1, \Sigma_1, \dots, \pi_k, \vec{\mu}_k, \Sigma_k)$

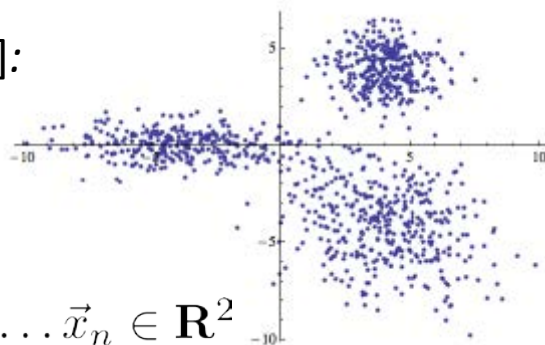
$$P[\vec{x} | \theta] = \sum_{i=1}^k \pi_i \frac{1}{\sqrt{(2\pi)^d \det(\Sigma_i)}} \exp \left\{ -\frac{1}{2} (\vec{x} - \vec{\mu}_i)^T \Sigma_i^{-1} (\vec{x} - \vec{\mu}_i) \right\}$$

Mixing weight

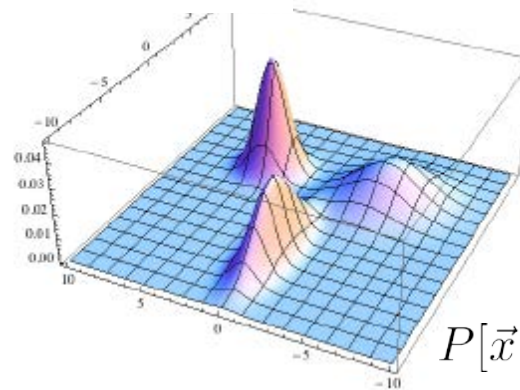
Mixture component

(this is called a mixture model)

Example in $\mathbf{R}^2 \times [3]$:



$\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \in \mathbf{R}^2$



$P[\vec{x} | \theta]$

GMM: Parameter Learning

$$P[\vec{x} \mid \theta] = \sum_{i=1}^k \pi_i \frac{1}{\sqrt{(2\pi)^d \det(\Sigma_i)}} \exp \left\{ -\frac{1}{2} (\vec{x} - \vec{\mu}_i)^\top \Sigma_i^{-1} (\vec{x} - \vec{\mu}_i) \right\}$$

$$\theta = (\pi_1, \vec{\mu}_1, \Sigma_1, \dots, \pi_k, \vec{\mu}_k, \Sigma_k)$$

So... how to learn the parameters θ ?

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MLE approach:

Given data $\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \in \mathbb{R}^d$

$$\theta_{\text{MLE}} := \arg \max_{\theta} \sum_{i=1}^n \ln P[\vec{x}_i \mid \theta]$$

$$= \arg \max_{\theta} \sum_{i=1}^n \ln \left[\sum_{j=1}^k \pi_j \frac{1}{\sqrt{(2\pi)^d \det(\Sigma_j)}} \exp \left\{ -\frac{1}{2} (\vec{x}_i - \vec{\mu}_j)^\top \Sigma_j^{-1} (\vec{x}_i - \vec{\mu}_j) \right\} \right]$$

ummm.... now what?

Cannot really simplify further!

GMM: Maximum Likelihood

MLE for Mixture modeling (like GMMs) is NOT a convex optimization problem

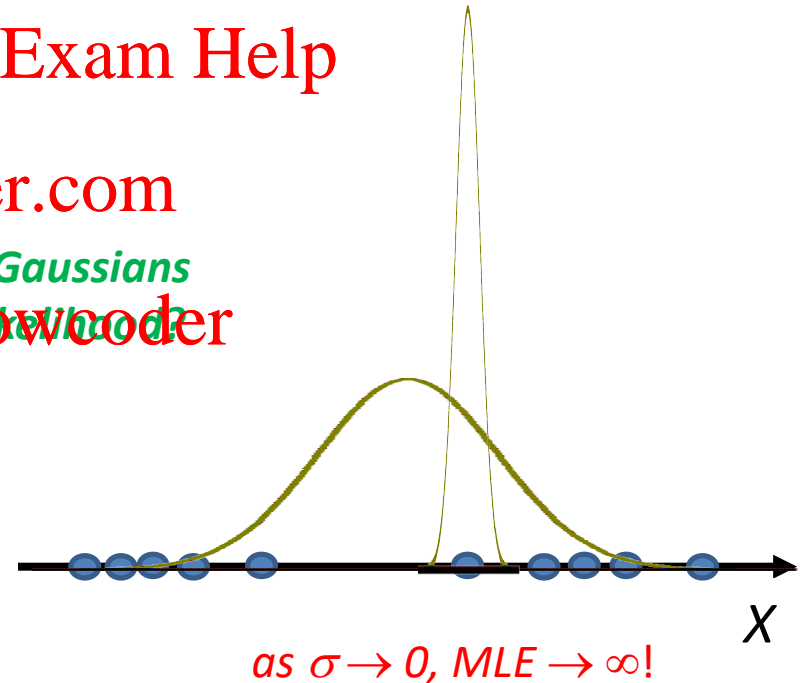
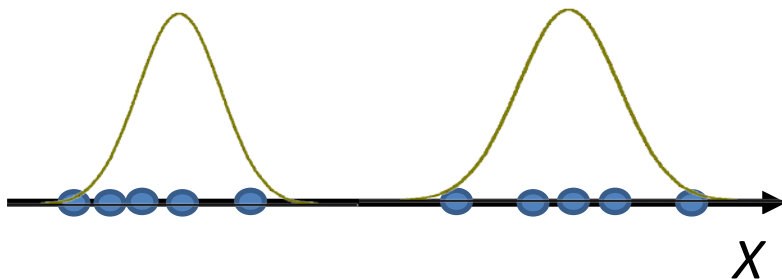
In fact **Maximum** Likelihood Estimate for GMMs is degenerate!

$X = \mathbf{R}$, $k = 2$ (fit two Gaussians in 1D):

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Which pair of Gaussians
gives higher likelihood?
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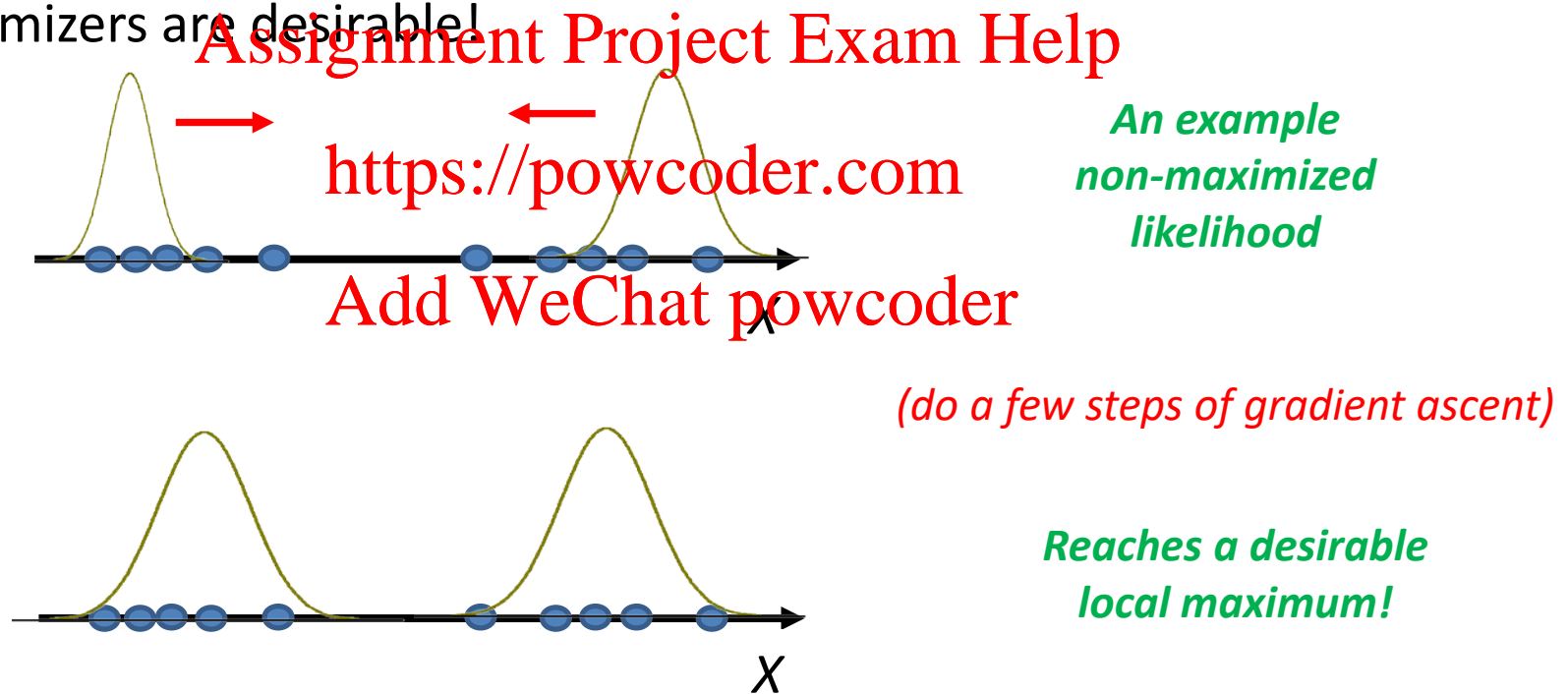


Aside: why doesn't this occur when fitting one Gaussian?

GMM: (local) Maximum Likelihood

So, can we make any progress?

Observation: even though a global MLE maximizer is not appropriate, several local maximizers are desirable!



A better algorithm for finding good parameters: Expectation Maximization (EM)

Expectation Maximization (EM) Algorithm

Similar in spirit to the alternating update for k -means algorithm

Idea:

- Initialize the parameters arbitrarily
- Given the current setting of parameters find the best (soft) assignment of data samples to the clusters (**Expectation-step**)
- Update all the parameters with respect to the current (soft) assignment that maximizes the likelihood (**Maximization-step**)
- Repeat until no more progress is made.

EM for GMM

Initialize $\theta = (\pi_1, \vec{\mu}_1, \Sigma_1, \dots, \pi_k, \vec{\mu}_k, \Sigma_k)$ arbitrarily

Expectation-step: For each $i \in \{1, \dots, n\}$ and $j \in \{1, \dots, k\}$ compute the assignment $w_j^{(i)}$ of data x_i to cluster j

$$w_j^{(i)} := \frac{\pi_j \sqrt{\det(\Sigma_j^{-1})} \exp\left(-\frac{1}{2}(\vec{x} - \vec{\mu}_j)^T \Sigma_j^{-1} (\vec{x} - \vec{\mu}_j)\right)}{\sum_{j'=1}^k \pi_{j'} \sqrt{\det(\Sigma_{j'}^{-1})} \exp\left(-\frac{1}{2}(\vec{x} - \vec{\mu}_{j'})^T \Sigma_{j'}^{-1} (\vec{x} - \vec{\mu}_{j'})\right)}$$

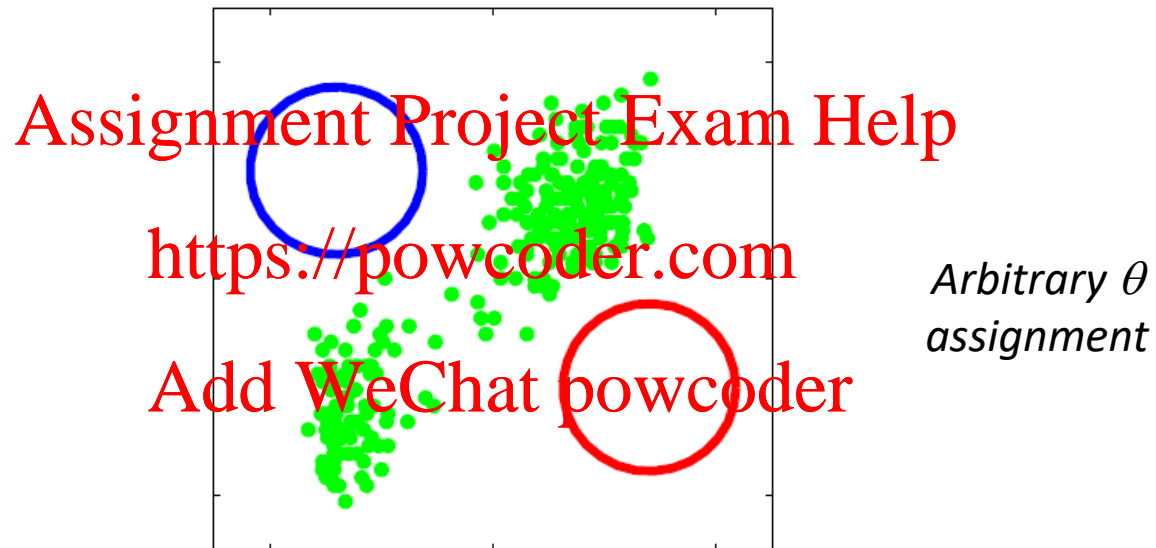
Maximization-step: Maximize the log-likelihood of the parameters

$$n_j := \sum_{i=1}^n w_j^{(i)} \quad \left(\begin{array}{l} \text{Effective number of points} \\ \text{assigned to cluster } j \end{array} \right) \quad \pi_j := \frac{n_j}{n}$$

$$\vec{\mu}_j := \frac{1}{n_j} \sum_{i=1}^n w_j^{(i)} \vec{x}_i \quad \Sigma_j := \frac{1}{n_j} \sum_{i=1}^n w_j^{(i)} (\vec{x}_i - \vec{\mu}_j)(\vec{x}_i - \vec{\mu}_j)^T$$

Why?

EM for GMM in Action

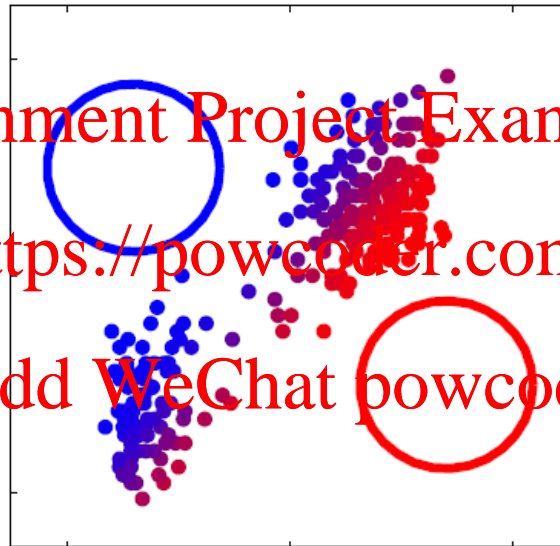


EM for GMM in Action

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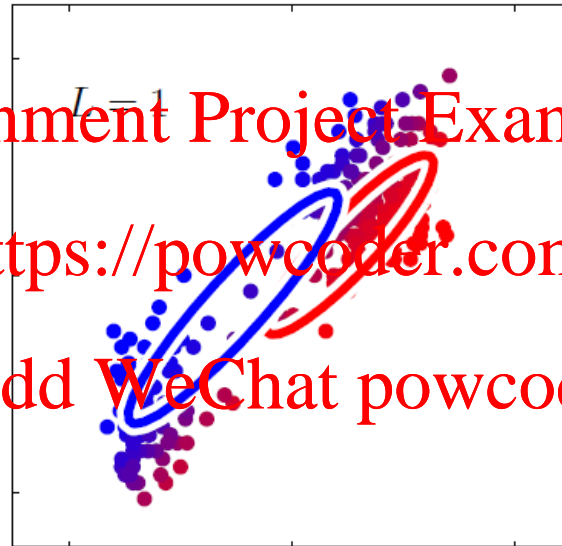
*E step: soft
assignment of data*

EM for GMM in Action

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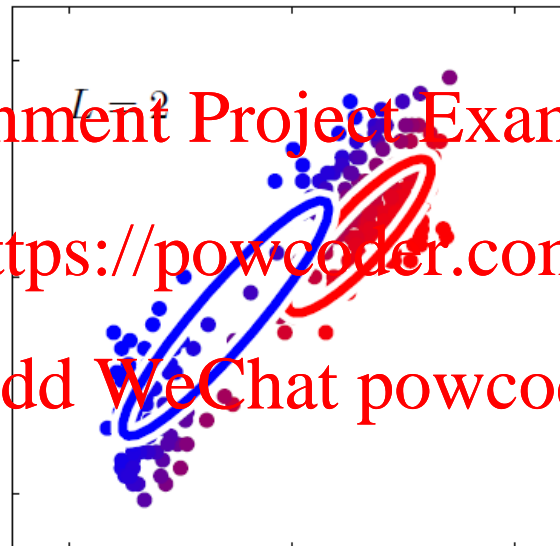
*M step: Maximize
parameter estimate*

EM for GMM in Action

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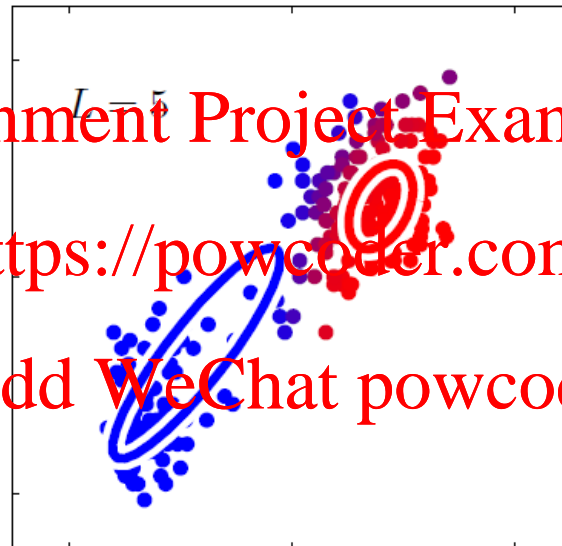
After two rounds

EM for GMM in Action

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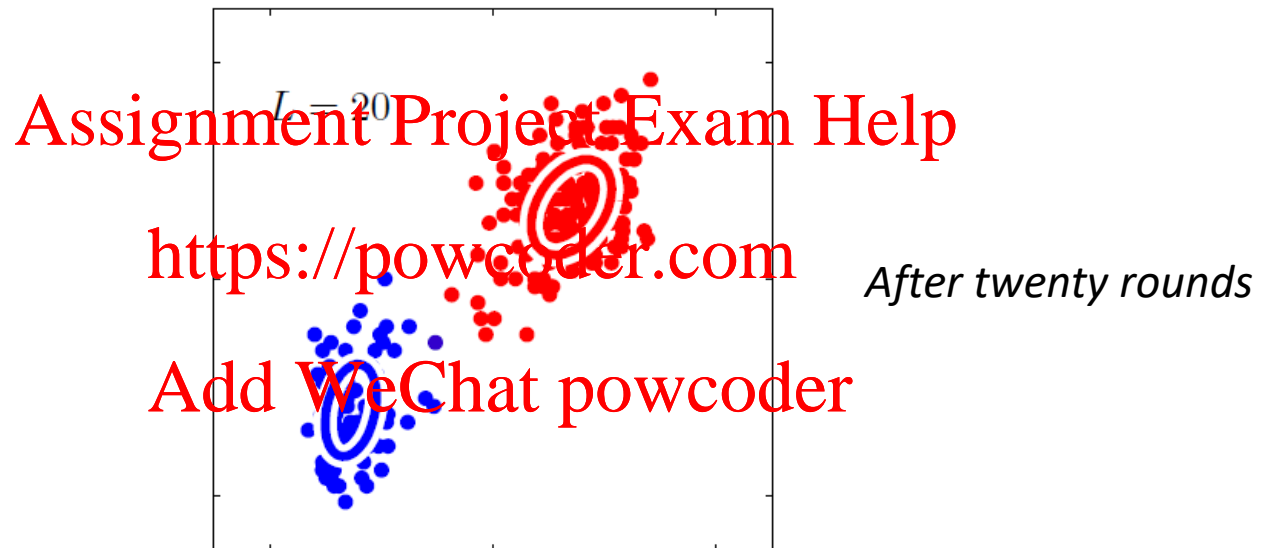
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After five rounds

EM for GMM in Action



What We Learned...

- Unsupervised Learning problems:
Clustering and Dimensionality Reduction
- K-means <https://powcoder.com>
- Hierarchical Clustering
- Gaussian Mixture Models
- EM algorithm

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Questions?

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Next time...

Dimension reduction! Assignment Project Exam Help

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