

# CSC 411 Lecture 15: K-Means

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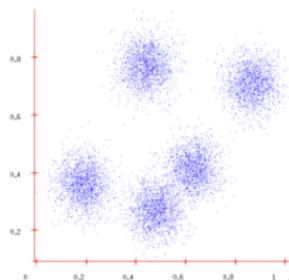
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# Motivating Examples

- Some examples of situations where you'd use unsupervised learning
  - ▶ You want to understand how a scientific field has changed over time. You want to take a large database of papers and model how the distribution of topics changes from year to year. But what are the topics?
  - ▶ You're a biologist studying animal behavior, so you want to infer a high-level description of their behavior from video. You don't know the set of behaviors ahead of time.
  - ▶ You want to reduce your energy consumption, so you take a time series of your energy consumption over time, and try to break it down into separate components (refrigerator, washing machine, etc.).
- Common theme: you have some data, and you want to infer the structure underlying the data.
- This structure is **latent**, which means it's never observed.

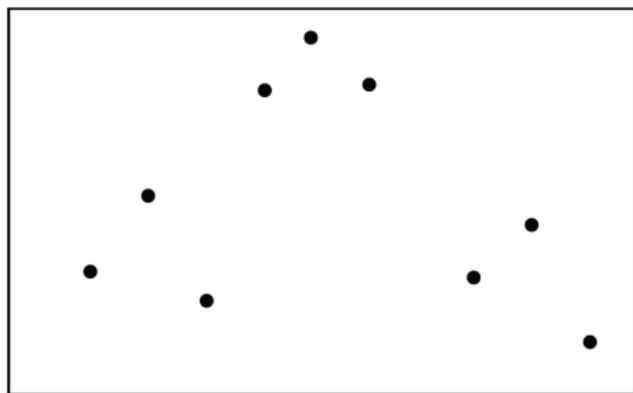
- In last lecture, we looked at density modeling where all the random variables were fully observed.
- The more interesting case is when some of the variables are latent, or never observed. These are called **latent variable models**.
  - ▶ Today's lecture: K-means, a simple algorithm for **clustering**, i.e. grouping data points into clusters
  - ▶ Next 2 lectures: reformulate clustering as a latent variable model, apply the EM algorithm

- Sometimes the data form clusters, where examples within a cluster are similar to each other, and examples in different clusters are dissimilar:



- Such a distribution is **multimodal**, since it has multiple **modes**, or regions of high probability mass.
- Grouping data points into clusters, with no labels, is called **clustering**
- E.g. clustering machine learning papers based on topic (deep learning, Bayesian models, etc.)

# Clustering



- Assume the data  $\{\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(N)}\}$  lives in a Euclidean space,  $\mathbf{x}^{(n)} \in \mathbb{R}^d$ .
- Assume each data point belongs to one of  $K$  clusters
- Assume the data points from same cluster are similar, i.e. close in Euclidean distance.
- How can we identify those clusters (data points that belong to each cluster)?

# K-means Objective

Let's formulate this as an optimization problem

- **K-means Objective:**

Find cluster centers  $\{\mathbf{m}_k\}_{k=1}^K$  and assignments  $\{r^{(n)}\}_{n=1}^N$  to minimize the sum of squared distances of data points  $\{\mathbf{x}^{(n)}\}$  to their assigned cluster centers

- Mathematically:

$$\min_{\{\mathbf{m}_k\}, \{r^{(n)}\}} J(\{\mathbf{m}_k\}, \{r^{(n)}\}) = \min_{\{\mathbf{m}\}, \{r\}} \sum_{n=1}^N \sum_{k=1}^K r_k^{(n)} \|\mathbf{m}_k - \mathbf{x}^{(n)}\|^2 \quad (1)$$

where  $r_k^{(n)} = \mathbb{I}[\mathbf{x}^{(n)} \text{ is assigned to cluster } k]$

- Finding an optimal solution is an NP-hard problem!

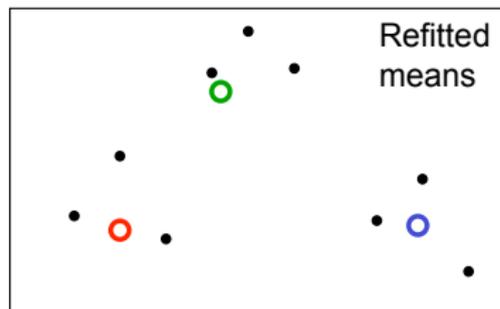
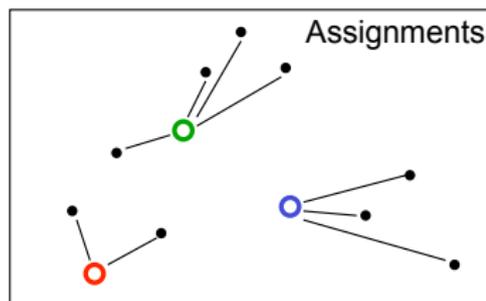
# Coordinate Descent

- But note:
  - ▶ If we fix the centers  $\{\mathbf{m}_k\}$  then we can easily find the optimal assignments  $\{\mathbf{r}^{(n)}\}$ 
    - ▶ Assign each point to the cluster with the nearest center (check!)
  - ▶ Likewise, if we fix the assignments  $\{\mathbf{r}^{(n)}\}$  then can easily find optimal centers  $\{\mathbf{m}_k\}$ 
    - ▶ Set each cluster's center to the average of its assigned data points (check!)
- Let's alternate between minimizing  $J(\{\mathbf{m}\}, \{\mathbf{r}\})$  with respect to  $\{\mathbf{m}\}$  and  $\{\mathbf{r}\}$
- This is called **coordinate descent**

# K-means

High level overview of algorithm:

- **Initialization:** randomly initialize cluster centers
- The algorithm iteratively alternates between two steps:
  - ▶ **Assignment step:** Assign each data point to the closest cluster
  - ▶ **Refitting step:** Move each cluster center to the mean of the data assigned to it



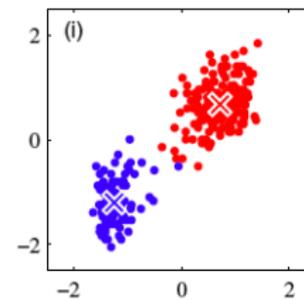
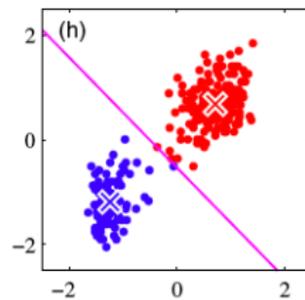
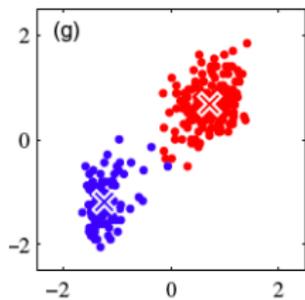
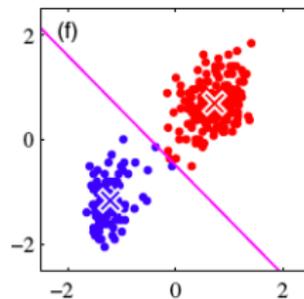
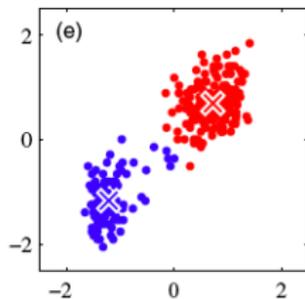
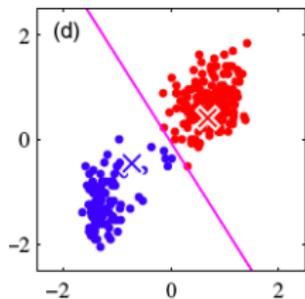
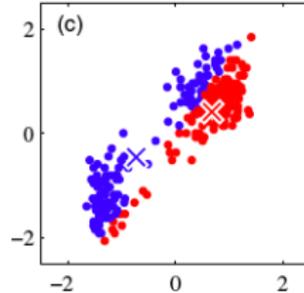
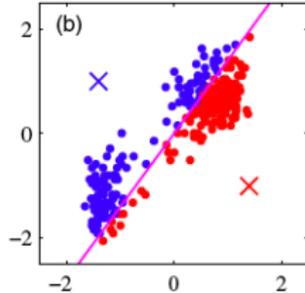
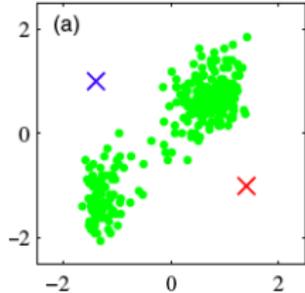


Figure from Bishop

Simple demo: <http://syskall.com/kmeans.js/>

# The K-means Algorithm

- **Initialization:** Set  $K$  cluster means  $\mathbf{m}_1, \dots, \mathbf{m}_K$  to random values
- Repeat until convergence (until assignments do not change):
  - ▶ **Assignment:** Optimize  $J$  w.r.t.  $\{\mathbf{r}\}$ : Each data point  $\mathbf{x}^{(n)}$  assigned to nearest center

$$\hat{k}^{(n)} = \arg \min_k \|\mathbf{m}_k - \mathbf{x}^{(n)}\|^2$$

and **Responsibilities** (1-hot encoding)

$$r_k^{(n)} = \mathbb{I}[\hat{k}^{(n)} = k]$$

- ▶ **Refitting:** Optimize  $J$  w.r.t.  $\{\mathbf{m}\}$ : Each center is set to mean of data assigned to it

$$\mathbf{m}_k = \frac{\sum_n r_k^{(n)} \mathbf{x}^{(n)}}{\sum_n r_k^{(n)}}$$

# K-means for Vector Quantization

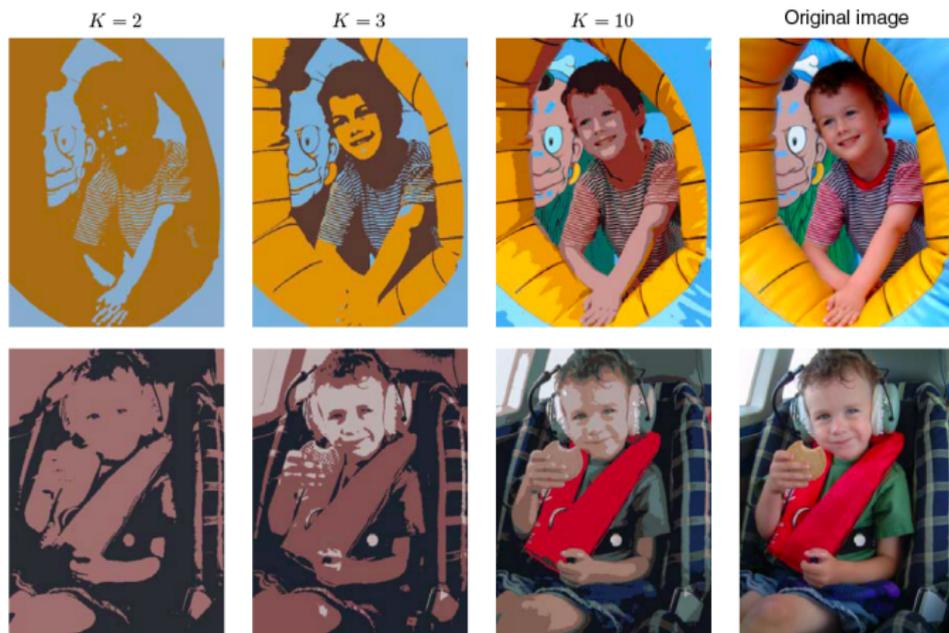
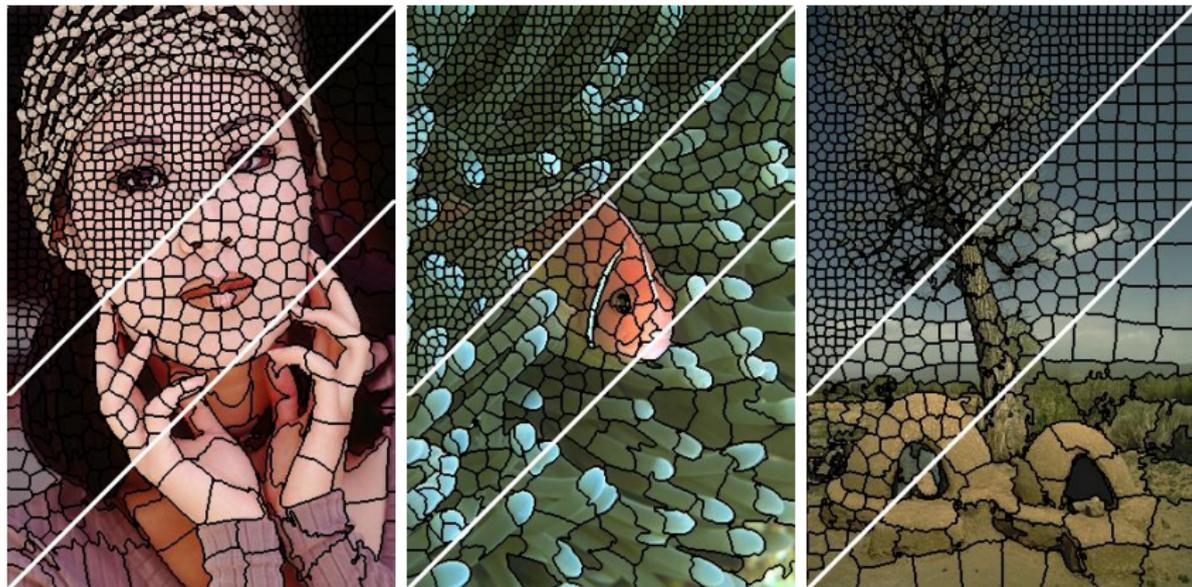


Figure from Bishop

- Given image, construct “dataset” of pixels represented by their RGB pixel intensities
- Run k-means, replace each pixel by its cluster center

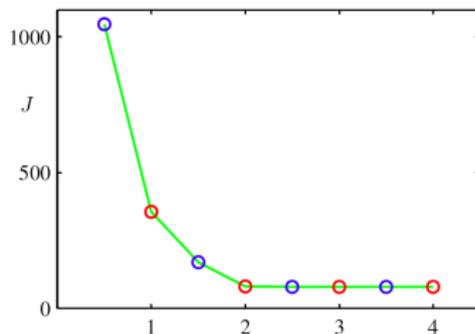
# K-means for Image Segmentation



- Given image, construct “dataset” of pixels, represented by their RGB pixel intensities and grid locations
- Run k-means (with some modifications) to get superpixels

# Why K-means Converges

- Whenever an assignment is changed, the sum squared distances  $J$  of data points from their assigned cluster centers is reduced.
- Whenever a cluster center is moved,  $J$  is reduced.
- **Test for convergence:** If the assignments do not change in the assignment step, we have converged (to at least a local minimum).
- This will always happen after a finite number of iterations, since the number of possible cluster assignments is finite

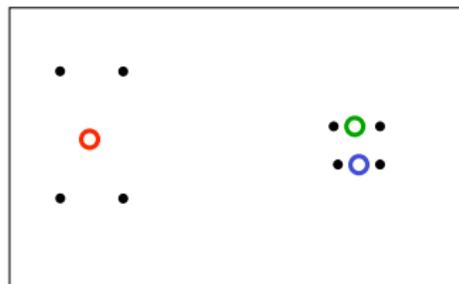


- K-means cost function after each assignment step (blue) and refitting step (red). The algorithm has converged after the third refitting step

# Local Minima

- The objective  $J$  is non-convex (so coordinate descent on  $J$  is not guaranteed to converge to the global minimum)
- There is nothing to prevent k-means getting stuck at local minima.
- We could try many random starting points
- We could try non-local split-and-merge moves:
  - ▶ Simultaneously **merge** two nearby clusters
  - ▶ and **split** a big cluster into two

A bad local optimum



- Instead of making hard assignments of data points to clusters, we can make **soft assignments**. One cluster may have a responsibility of .7 for a datapoint and another may have a responsibility of .3.
  - ▶ Allows a cluster to use more information about the data in the refitting step.
  - ▶ How do we decide on the soft assignments?

# Soft K-means Algorithm

- **Initialization:** Set  $K$  means  $\{\mathbf{m}_k\}$  to random values
- Repeat until convergence (measured by how much  $J$  changes):
  - ▶ **Assignment:** Each data point  $n$  given soft "degree of assignment" to each cluster mean  $k$ , based on responsibilities

$$r_k^{(n)} = \frac{\exp[-\beta d(\mathbf{m}_k, \mathbf{x}^{(n)})]}{\sum_j \exp[-\beta d(\mathbf{m}_j, \mathbf{x}^{(n)})]}$$

$$\implies \mathbf{r}^{(n)} = \text{softmax}(-\beta[d(\mathbf{m}_k, \mathbf{x}^{(n)})]_{k=1}^K)$$

- ▶ **Refitting:** Model parameters, means, are adjusted to match sample means of datapoints they are responsible for:

$$\mathbf{m}_k = \frac{\sum_n r_k^{(n)} \mathbf{x}^{(n)}}{\sum_n r_k^{(n)}}$$

# Questions about Soft K-means

Some remaining issues

- How to set  $\beta$ ?
- Clusters with unequal weight and width?

These aren't straightforward to address with K-means. Instead, next lecture, we'll reformulate clustering using a generative model.