Lecture 11

Clustering: K-means and Mean Shift

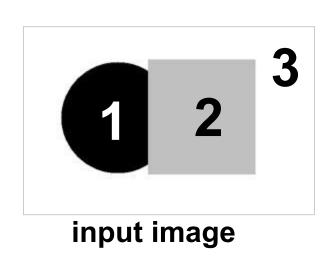
What will we learn today?

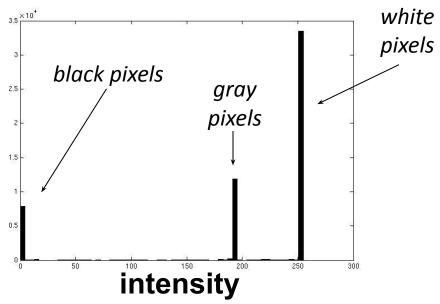
- K-means clustering
- Mean-shift clustering

Reading: Szeliski Chapters: 5.2.2, 7.5.2

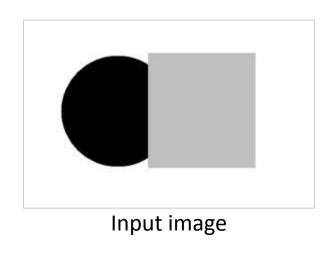
D. Comaniciu and P. Meer, Mean Shift: A Robust Approach toward Feature Space Analysis, PAMI 2002.

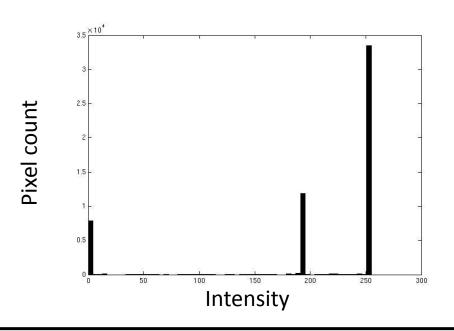
Image Segmentation: Toy Example

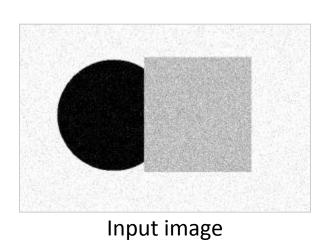




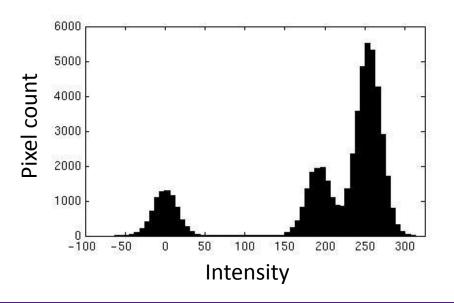
- These intensities define the three groups.
- We could label every pixel in the image according to which of these primary intensities it is.
 - o i.e., segment the image based on the intensity feature.
- What if the image isn't quite so simple?

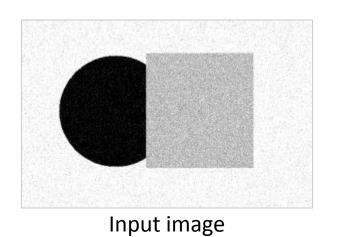


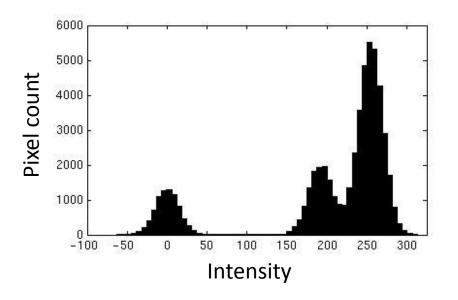




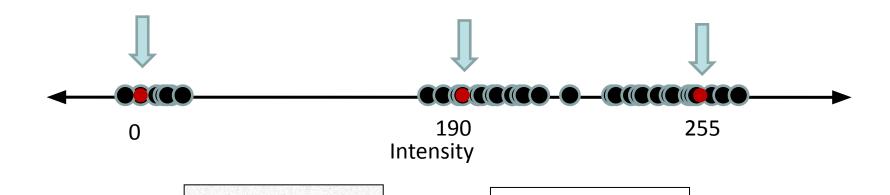
Slide credit: Kristen Grauman

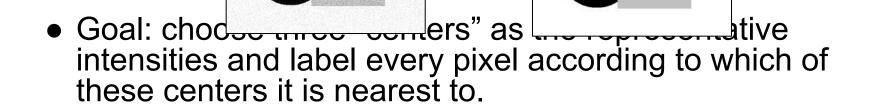






- Now how to determine the three main intensities that define our groups?
- We need to cluster.





 Best cluster centers are those that minimize Sum of Square Distance (SSD) between all points and their nearest cluster center c_i:

$$SSD = \sum_{clusteri} \sum_{x \in clusteri} (x - c_i)^2$$

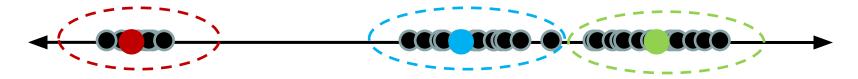
Clustering

Goal: cluster to minimize variance in data given clusters

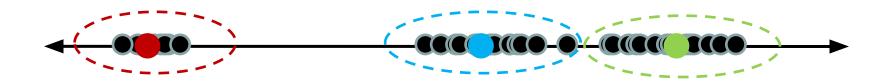
Cluster center Data
$$c^*, \ \delta^* = \underset{c, \, \delta}{\operatorname{arg\,min}} \frac{1}{N} \sum_{j=1}^{N} \sum_{i}^{K} \delta_{ij} \left(c_i - x_j \right)^2$$
 Whether x_j is assigned to c_i

Clustering

- With this objective, it is a "chicken and egg" problem:
 - o If we knew the *cluster centers*, we could allocate points to groups by assigning each to its closest center.



o If we knew the *group memberships*, we could get the centers by computing the mean per group.



- 1. Initialize $\ell = 0$): cluster centers,..., c_K
- 2. Compute δ^t : assign each point to the closest center δ^t denotes the set of assignment for each to cluster at iteration t

$$\delta^{t} = \underset{\delta}{\operatorname{argmin}} \frac{1}{N} \sum_{j}^{N} \sum_{i}^{K} \delta_{ij}^{t-1} \left(c_{i}^{t-1} - x_{j} \right)^{2}$$

3. Computer c^t : update cluster centers as the mean of the points

$$c^{t} = \underset{c}{\operatorname{argmin}} \frac{1}{N} \sum_{j}^{N} \sum_{i}^{K} \delta_{ij}^{t} \left(c_{i}^{t-1} x_{j} \right)^{2}$$

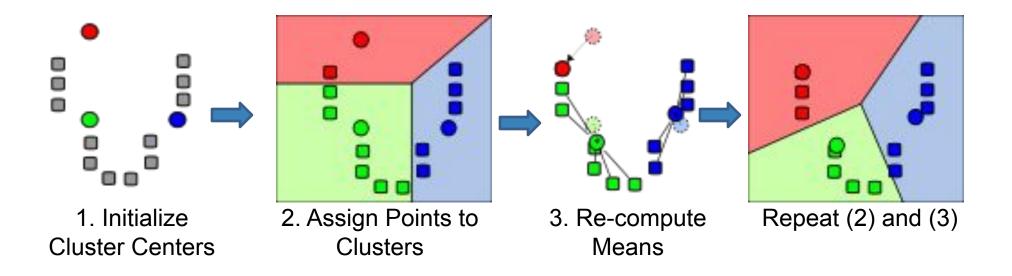
4. Update t = t + 1, Repeat Step 2-3 till stopped

- Initialize $\ell=0$): cluster centers,..., c_K Commonly used: random initialization

 - Or greedily choose K to minimize residual
- Compute S^t : assign each point to the closest center
 - Typical distance measure:
 - Euclidean $sim(x, x') = x^T x'$
 - Cosine $sim(x, x') = x^T x' / (||x|| \cdot ||x'||)$
 - Others
- Computer C^t : update cluster centers as the mean of the points

$$c^{t} = \underset{c}{\operatorname{argmin}} \frac{1}{N} \sum_{j}^{N} \sum_{i}^{K} \delta_{ij}^{t} \left(c_{i}^{t-1} x_{j} \right)^{2}$$

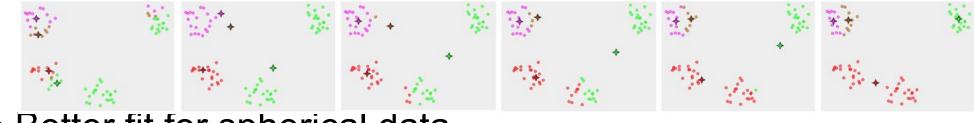
- 4. Update t = t + 1• C doesn't change anymore.



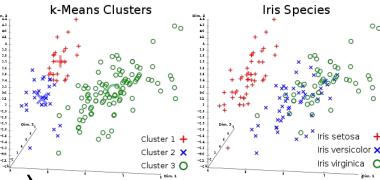
• Online demo:

https://kmeans.js.org/en/

- Converges to a local minimum solution
 - Initialize multiple runs



Better fit for spherical data



Need to pick K (# of clusters)

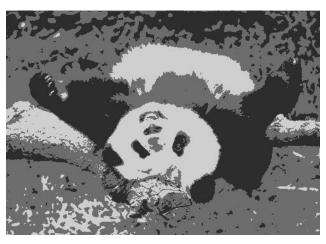
Segmentation as Clustering



Original image



2 clusters



K-Means++

- Can we prevent arbitrarily bad local minima?
- 1. Randomly choose first center.
- 2. Pick new center with prob. proportional $(x-c_i)^2$ o (Contribution of x to total error)
- 3. Repeat until *K* centers.
- Expected error_{= $O(\log K)$} * optimal

S<mark>l</mark>ide credit: Steve Seitz

<u> Arthur & Vassilvitskii 2007</u>

Feature Space

- Feature space: what measurements do we include in x_i ?
- Depending on what we choose as the feature space, we can group pixels in different ways.
- Grouping pixels based on intensity similarity



Feature space: intensity value (1)

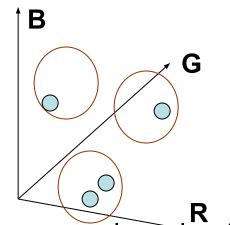


Feature Space

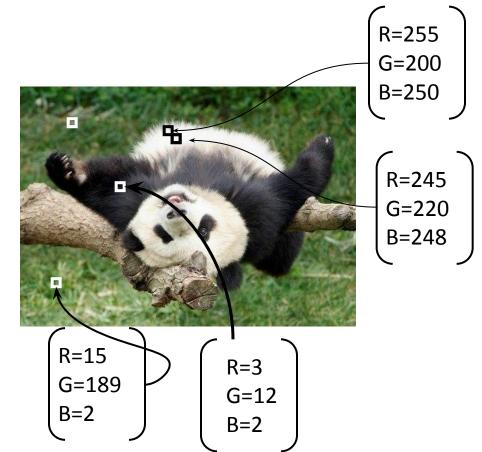
• Depending on what we choose as the *feature space*, we can group

pixels in different ways.

 Grouping pixels based on color similarity

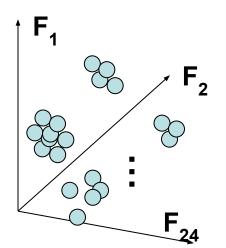


• Feature space: color value (3D)

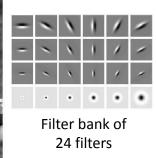


Feature Space

- Depending on what we choose as the feature space, we can group pixels in different ways.
- Grouping pixels based on texture similarity



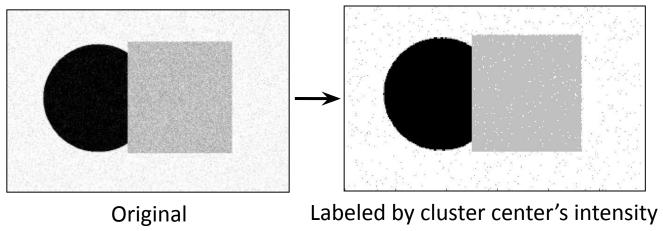




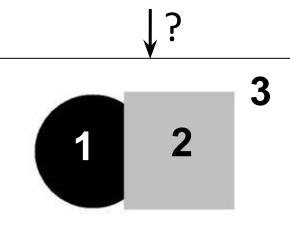
• Feature space: filter bank responses (e.g., 24D)

Smoothing Out Cluster Assignments

Assigning a cluster label per pixel may yield outliers:

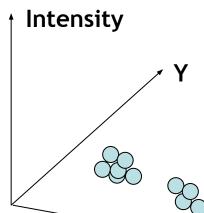


 How can we ensure they are spatially smooth?

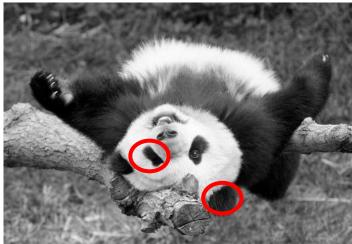


Segmentation as Clustering

- Depending on what we choose as the feature space, we can group pixels in different ways.
- Grouping pixels based on intensity+position similarity



⇒ Way to encode both similarity and proximity.



K-Means Clustering Results

- K-means clustering based on intensity or color is essentially vector quantization of the image attributes
 - Clusters don't have to be spatially coherent



Intensity-based clusters



Color-based clusters



nage source: Forsyth & P

K-Means Clustering Results

- K-means clustering based on intensity or color is essentially vector quantization of the image attributes
 - Clusters don't have to be spatially coherent
- Clustering based on (r,g,b,x,y) values enforces more spatial coherence



Image source: Forsyth & Ponce

How to evaluate clusters?

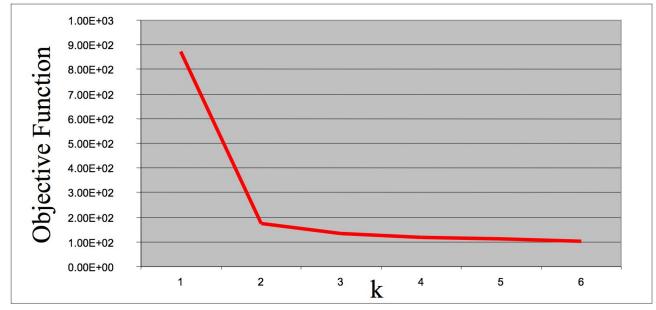
- Generative
 - Our How well are points reconstructed from the clusters?
- Discriminative
 - Our How well do the clusters correspond to labels?
 - Can we correctly classify which pixels belong to the panda?
 - Note: unsupervised clustering does not aim to be discriminative as we don't have the labels.

How to choose the number of clusters?

Try different numbers of clusters in a validation set and look at performance.

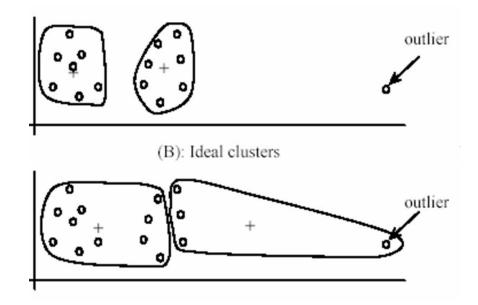
We can plot the objective function values for k equals 1 to 6...

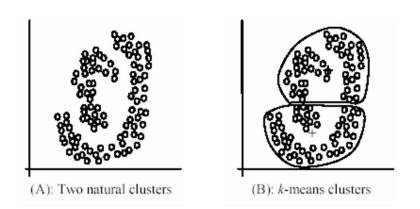
The abrupt change at k = 2, is highly suggestive of two clusters in the data. This technique for determining the number of clusters is known as "knee finding" or "elbow finding".



K-Means pros and cons

- Pros
 - Finds cluster centers that minimize conditional variance (good representation of data)
 - Simple and fast, Easy to implement
- Cons
 - Need to choose K
 - Sensitive to outliers
 - Prone to local minima
 - All clusters have the same parameters (e.g., distance measure is non-adaptive)
 - *Can be slow: each iteration is O(KNd) for N d-dimensional points
- Usage
 - Unsupervised clustering
 - Rarely used for pixel segmentation



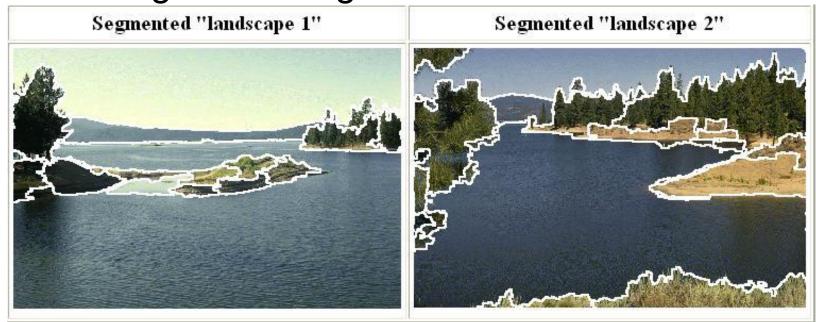


What will we learn today?

- K-means clustering
- Mean-shift clustering

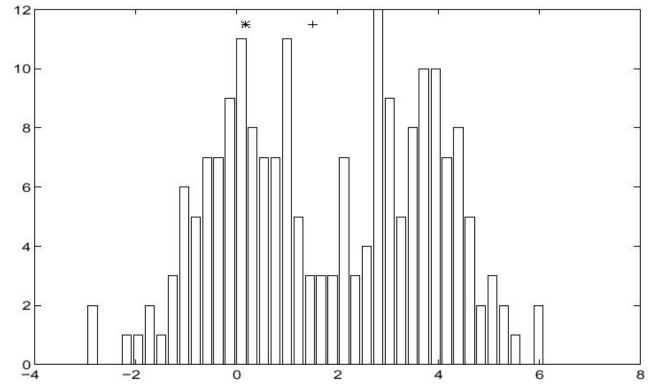
Mean-Shift Segmentation

 An advanced and versatile technique for clustering-based segmentation



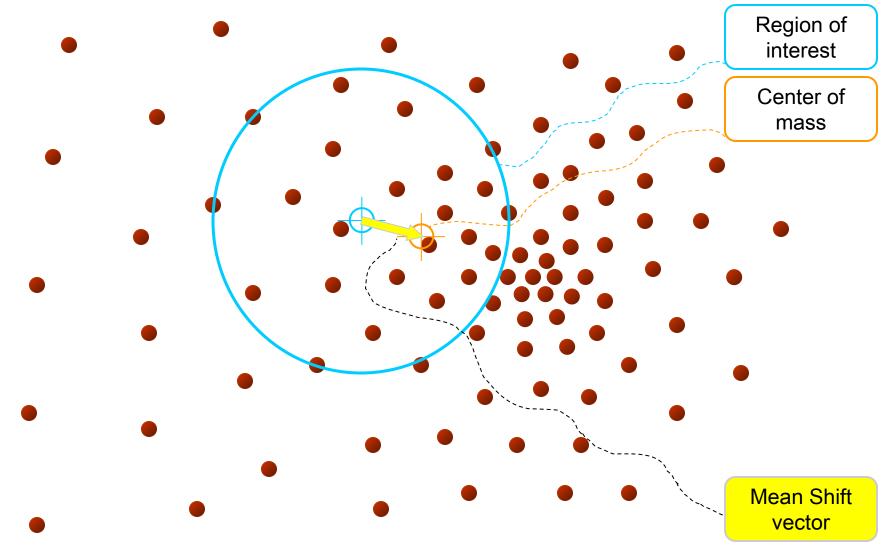
http://www.caip.rutgers.edu/~comanici/MSPAMI/msPamiResults.html

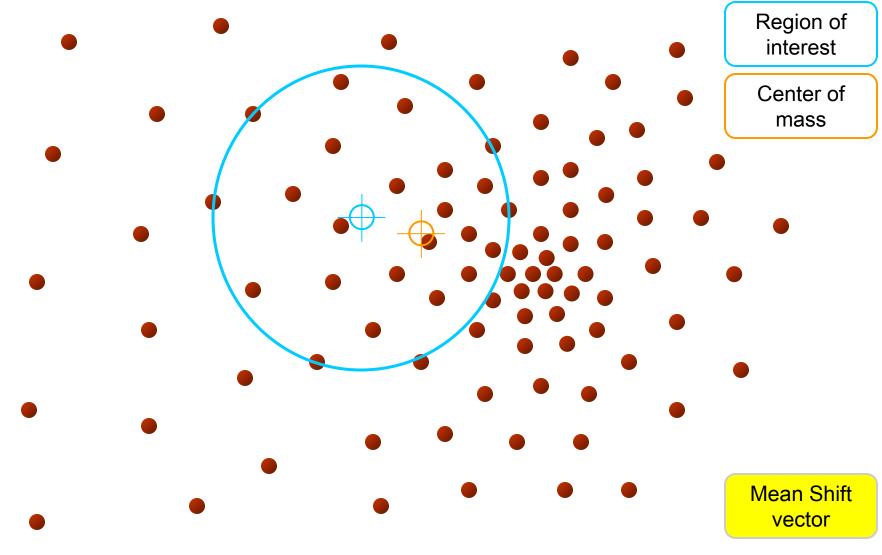
Mean-Shift Algorithm

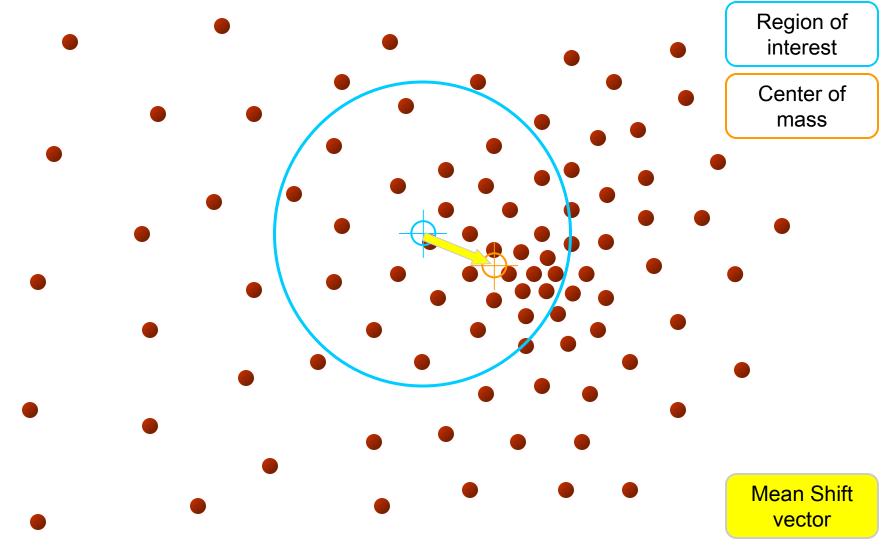


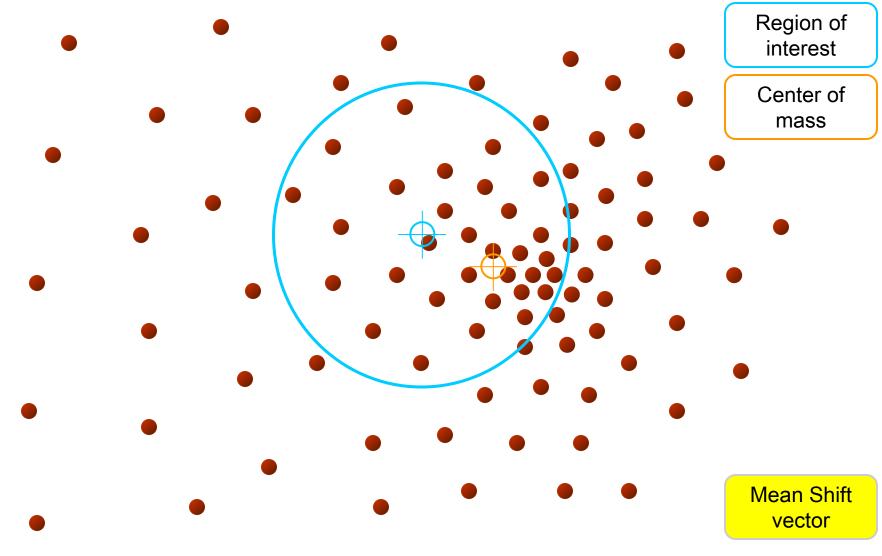
- Iterative Mode Se
 - 1. Initialize random seed, and window VV
 - 2. Calculate center of gravity (the "mean") of W:
 - 3. Shift the search window to the mean
 - 4. Repeat Step 2 until convergence

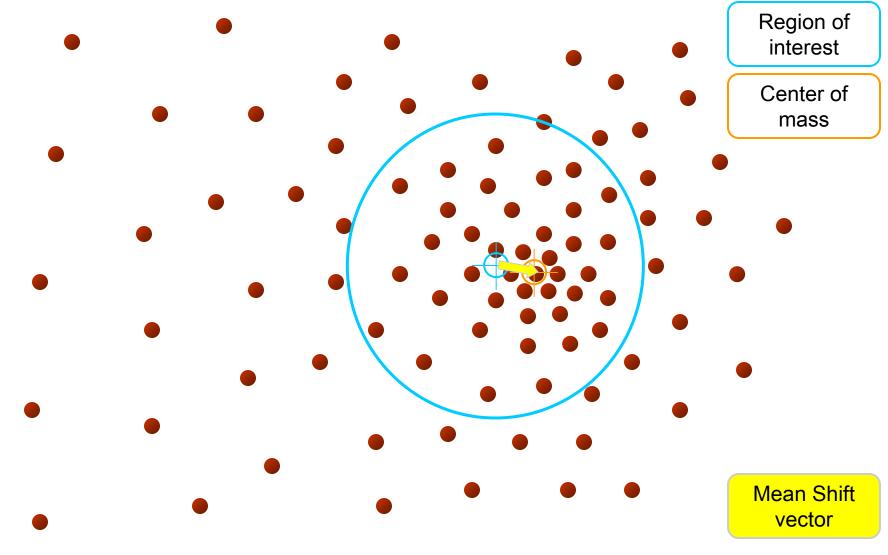
$$\sum_{x \in W} x H(x)$$

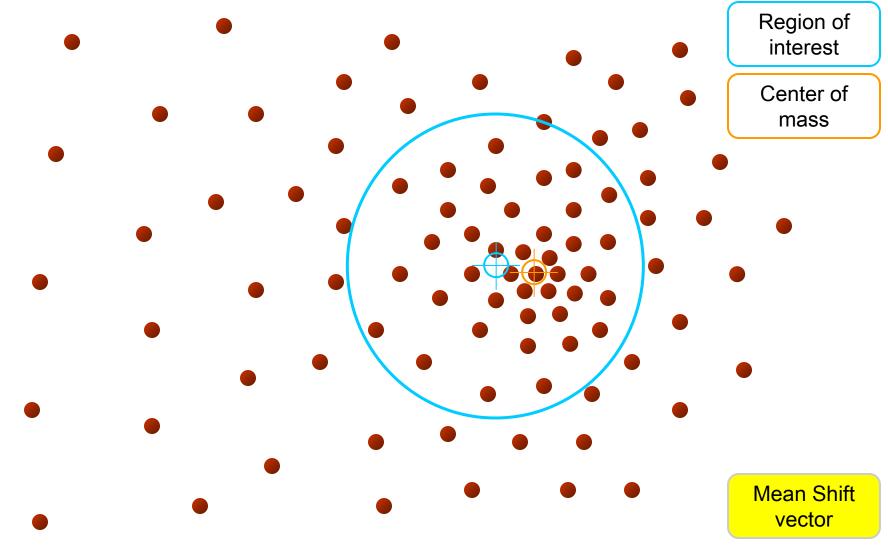


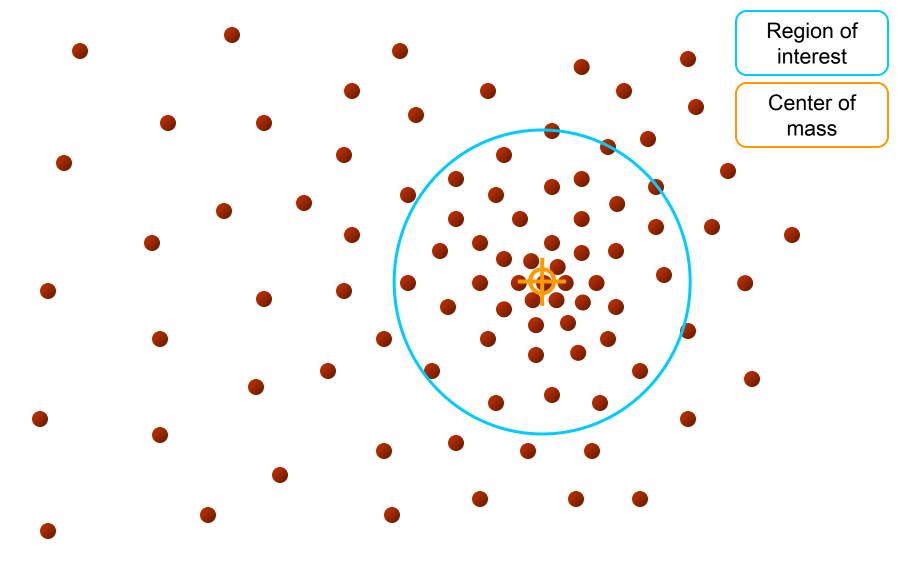




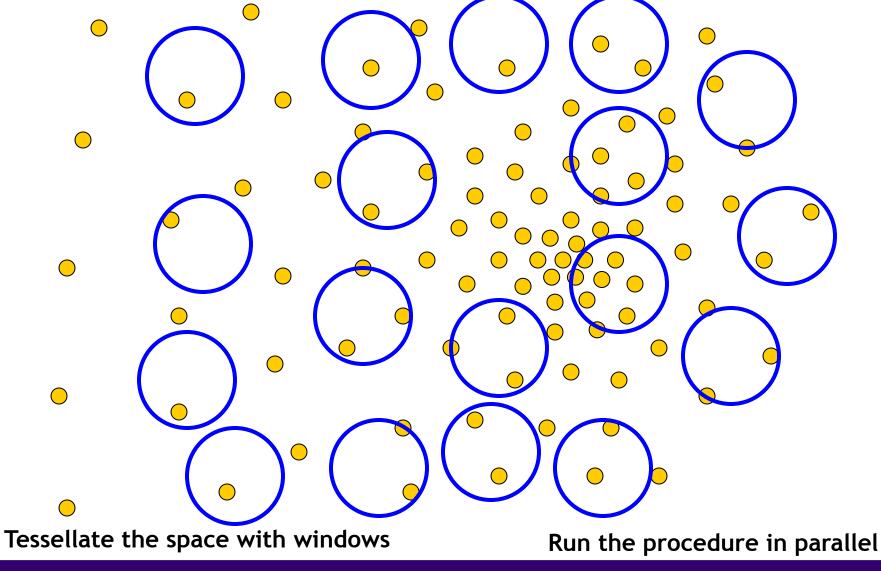




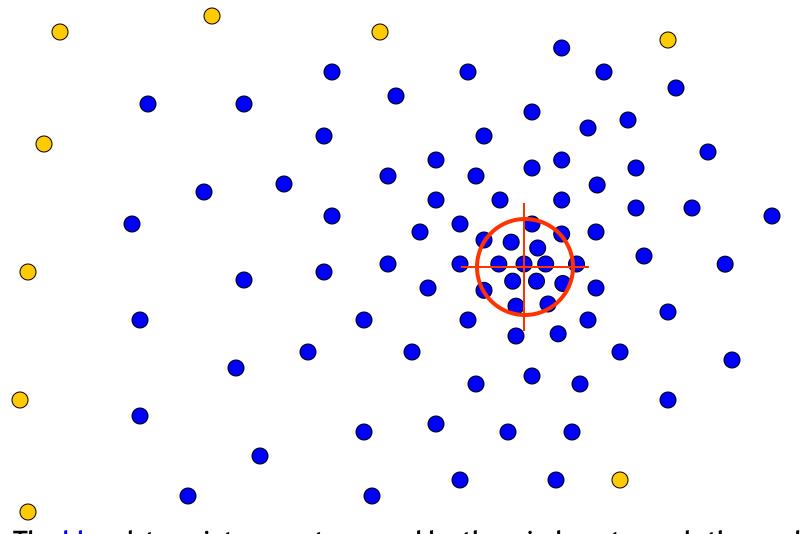




Real Modality Analysis



Real Modality Analysis

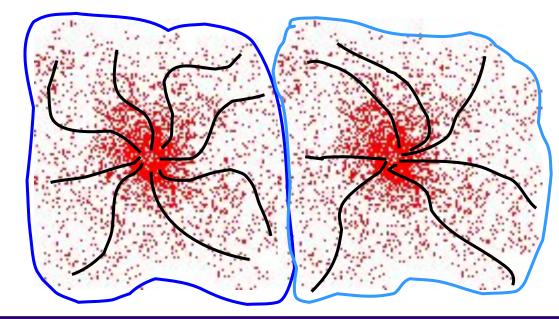


Slide by Y. Ukrainitz

The blue data points were traversed by the windows towards the mode.

Mean-Shift Clustering

- Cluster: all data points in the attraction basin of a mode
- Attraction basin: the region for which all trajectories lead to the same mode

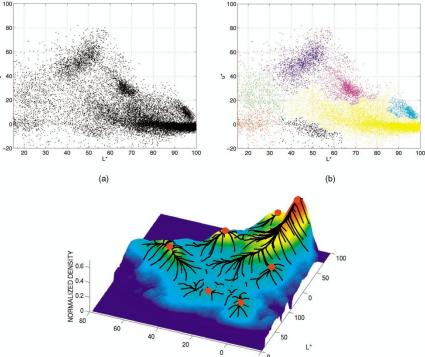


Mean-Shift Clustering/Segmentation

- Find features (color, gradients, texture, etc)
- Initialize windows at individual pixel locations
- Perform mean shift for each window until convergence

Merge windows that end up near the same "peak" or

mode



Mean-Shift Segmentation Results









ke credit: Svetlana Lazebnik







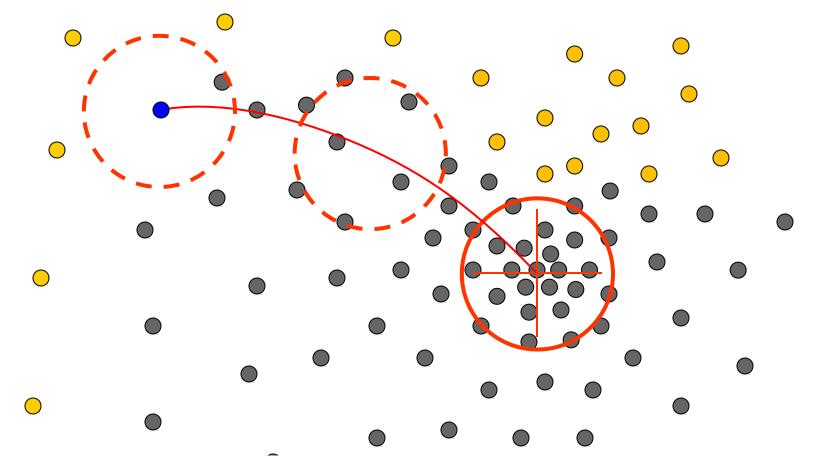


Moi

Ranjay Kri

Aph 90712024

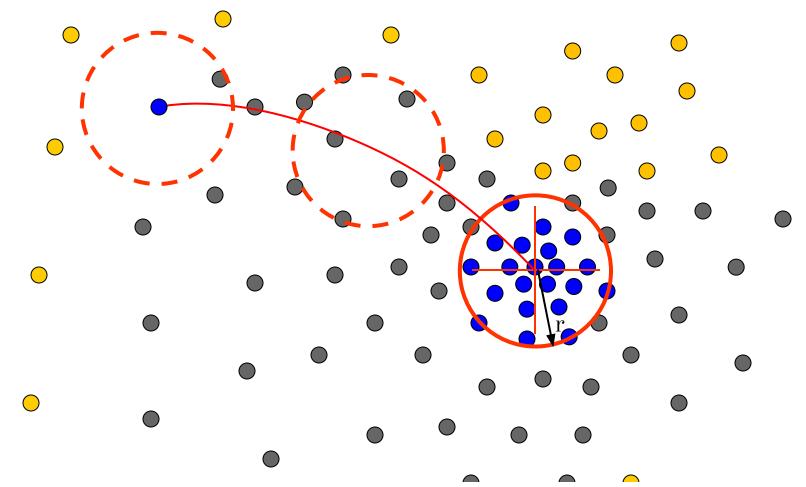
Problem: Computational Complexity



- Need to shift many windows...
- Many computations will be redundant.

Slide credit: Bastian Leibe

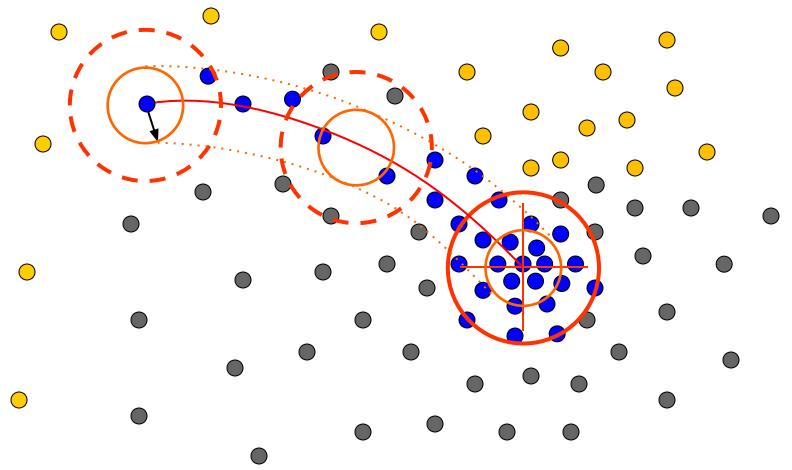
Speedups: Basin of Attraction



1. Assign all points within radius r of end point to the mode.

Slide credit: Bastian Leibe

Speedups



2. Assign all points within radius r/c of the search path to the mode -> reduce the number of data points to search.

Slide credit: Bastian Leibe

Technical Details

Given n data points $\mathbf{x}_i \in \mathbb{R}^d$, the multivariate kernel density estimate using a radially symmetric kernel¹ (e.g., Epanechnikov and Gaussian kernels), $K(\mathbf{x})$, is given by,

$$\hat{f}_K = \frac{1}{nh^d} \sum_{i=1}^n K\left(\frac{\mathbf{x} - \mathbf{x}_i}{h}\right),\tag{1}$$

where h (termed the bandwidth parameter) defines the radius of kernel. The radially symmetric kernel is defined as,

$$K(\mathbf{x}) = c_k k(\|\mathbf{x}\|^2), \tag{2}$$

where c_k represents a normalization constant.

Other Kernels

A kernel is a function that satisfies the following requirements :

$$1. \int_{R^d} \phi(x) = 1$$

2.
$$\phi(x) \ge 0$$

Some examples of kernels include:

1. Rectangular
$$\phi(x) = \begin{cases} 1 & a \leq x \leq b \\ 0 & else \end{cases}$$

2. Gaussian
$$\phi(x)=e^{-\frac{x^2}{2\sigma^2}}$$

3. Epanechnikov
$$\phi(x)= \begin{cases} \frac{3}{4}(1-x^2) & if \ |x|\leq 1 \\ 0 & else \end{cases}$$

<u>source</u>

Technical Details

Taking the derivative of:
$$\hat{f}_K = \frac{1}{nh^d} \sum_{i=1}^n K\left(\frac{\mathbf{x} - \mathbf{x}_i}{h}\right)$$

$$\nabla \hat{f}(\mathbf{x}) = \underbrace{\frac{2c_{k,d}}{nh^{d+2}} \left[\sum_{i=1}^{n} g\left(\left\| \frac{\mathbf{x} - \mathbf{x}_{i}}{h} \right\|^{2} \right) \right]}_{\text{term 1}} \underbrace{\left[\sum_{i=1}^{n} \mathbf{x}_{i} g\left(\left\| \frac{\mathbf{x} - \mathbf{x}_{i}}{h} \right\|^{2} \right) - \mathbf{x} \right]}_{\text{term 2}}, \tag{3}$$

where g(x) = -k'(x) denotes the derivative of the selected kernel profile.

- Term1: this is proportional to the density estimate at x (similar to equation 1 from two slides ago).
- Term2: this is the mean-shift vector that points towards the direction of maximum density.

Comaniciu & Meer, 2002

Technical Details

Finally, the mean shift procedure from a given point x_t is:

1. Compute the mean shift vector **m**:

$$\left[\frac{\sum_{i=1}^{n} \mathbf{x}_{i} g\left(\left\|\frac{\mathbf{x}-\mathbf{x}_{i}}{h}\right\|^{2}\right)}{\sum_{i=1}^{n} g\left(\left\|\frac{\mathbf{x}-\mathbf{x}_{i}}{h}\right\|^{2}\right)} - \mathbf{x}\right]$$

2. Translate the density window:

$$\mathbf{x}_i^{t+1} = \mathbf{x}_i^t + \mathbf{m}(\mathbf{x}_i^t).$$

3. Iterate steps 1 and 2 until convergence.

$$\nabla f(\mathbf{x}_i) = 0.$$

Comaniciu & Meer, 2002

Mean-Shift pros and cons

• Pros

- General, application-independent tool
- Model-free, does not assume any prior shape (spherical, elliptical, etc.) on data clusters
- Just a single parameter (window size h)
 - h has a physical meaning (unlike k-means)
- Finds variable number of modes
- Robust to outliers

Cons

- Output depends on window size
- Window size (bandwidth) selection is not trivial
- Computationally (relatively) expensive (~2s/image)
- Does not scale well with dimension of feature space

Summary

- K-means clustering
- Mean-shift clustering