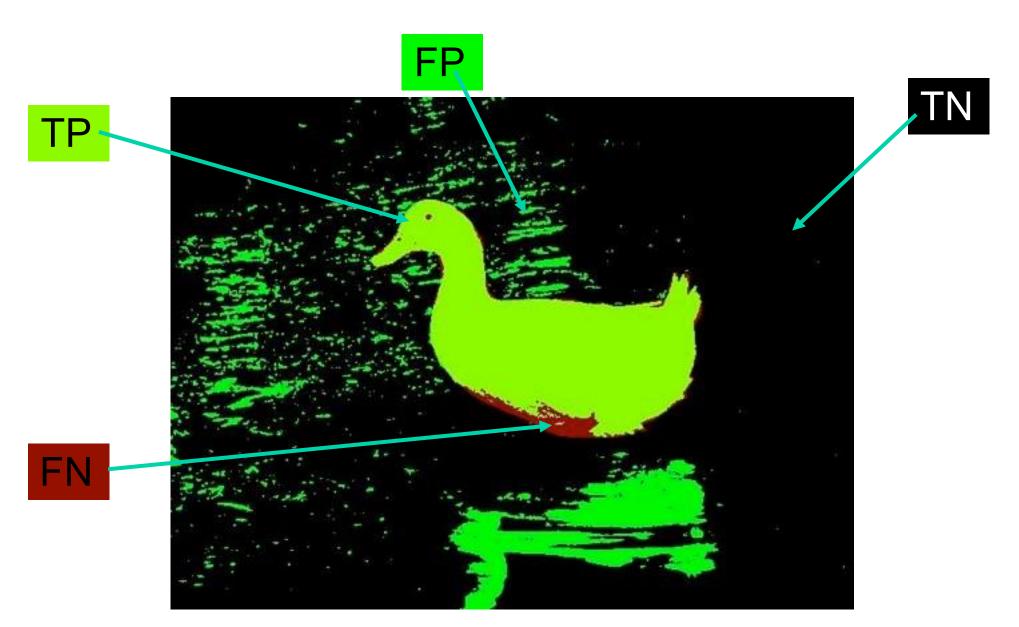


Classification outcomes



Limitations of Thresholding

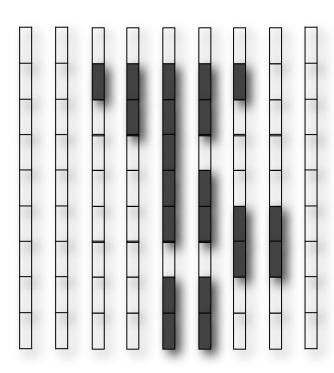
 Why can we segment images much better by eye than through thresholding processes?

 We might improve results by considering image context: Surface Coherence

Pixel Connectivity

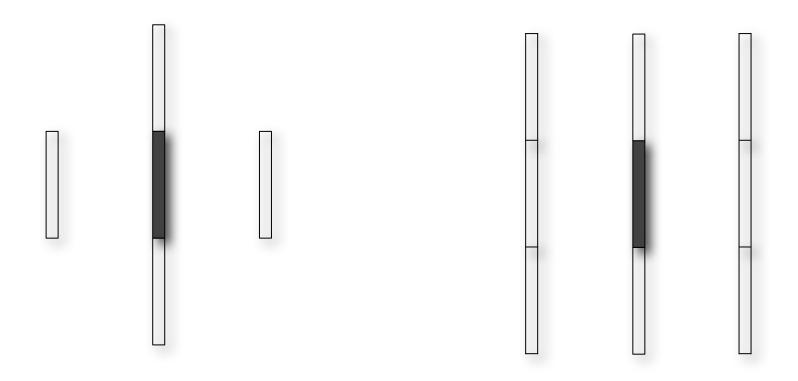
 We need to define which pixels are neighbours.

 Are the dark pixels in this array connected?



Pixel Neighborhoods

4-neighbourhood



8-neighbourhood

Pixel Paths

• A 4-connected path between pixels p_1 and p_n is a set of pixels $\{p_1, p_2, ..., p_n\}$ such that p_i is a 4-neighbour of p_{i+1} , i=1,...,n-1.

• In an 8-connected path, p_i is an 8-neighbour of p_{i+1} .

Connected Regions

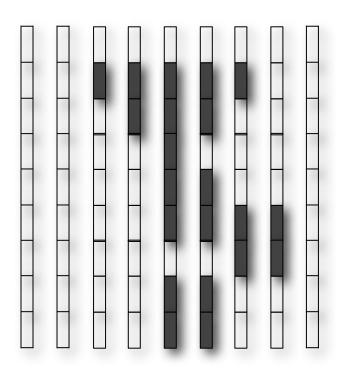
 A region is 4-connected if it contains a 4connected path between every pair of its pixels.

 A region is 8-connected if it contains an 8connected path between every pair of its pixels.

Connected Regions

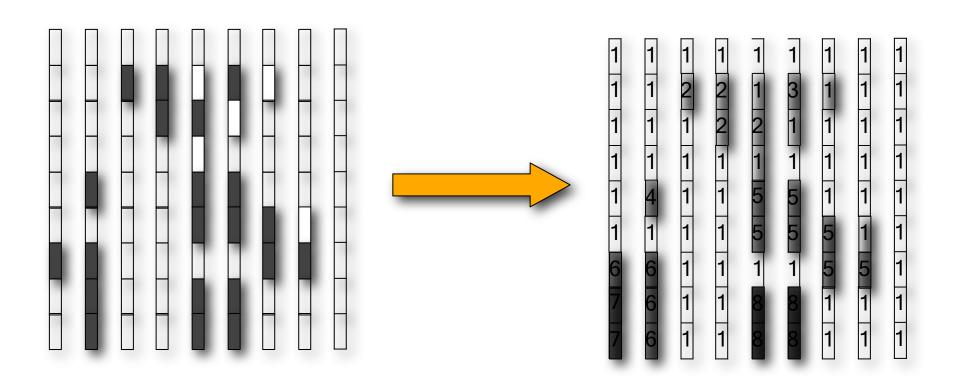
 Now what can we say about the dark pixels in this array?

What about the light pixels?



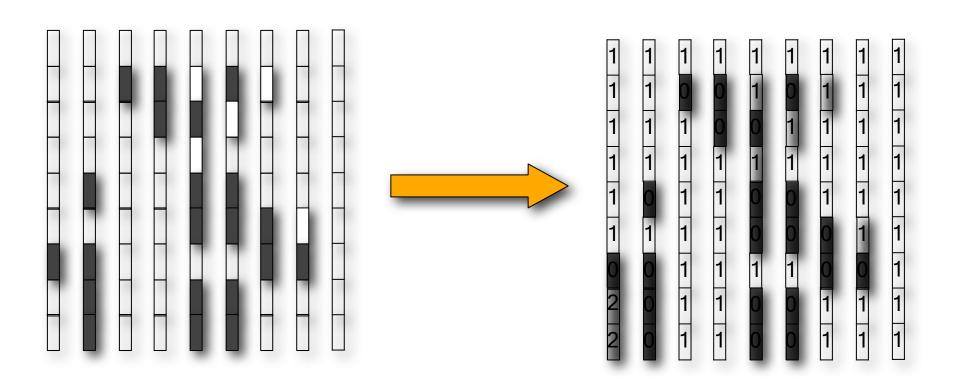
Connected Components Labelling

 Labels each connected component of a binary image with a separate number.



Foreground Labelling

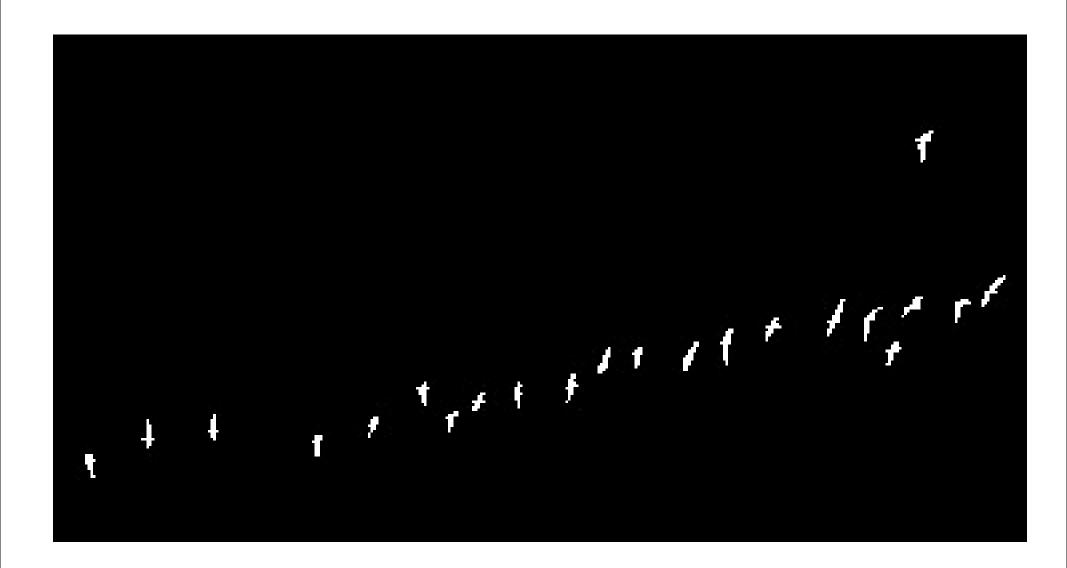
 Only extract the connected components of the foreground



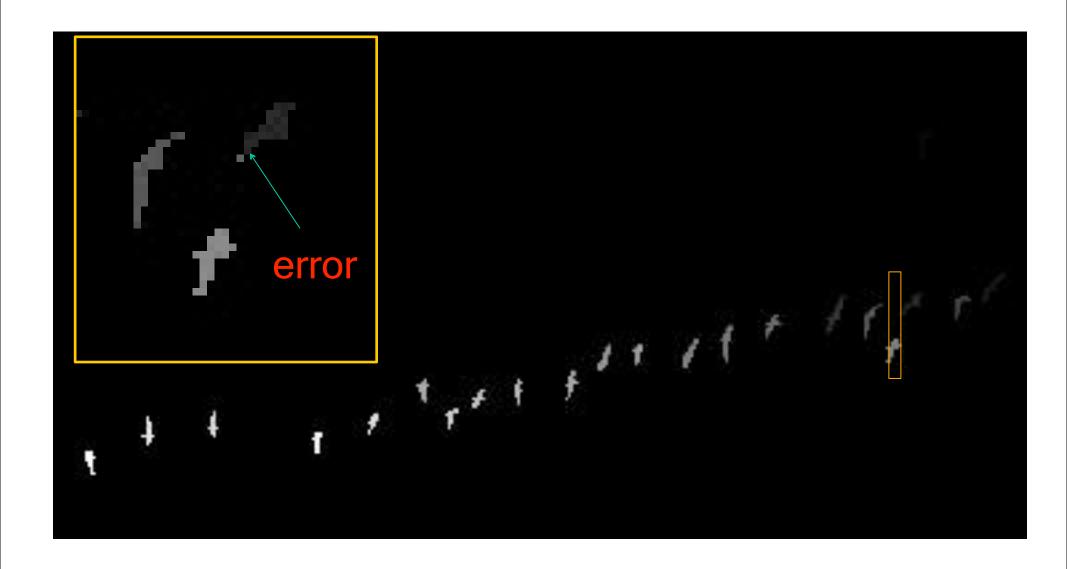
Goose Detector



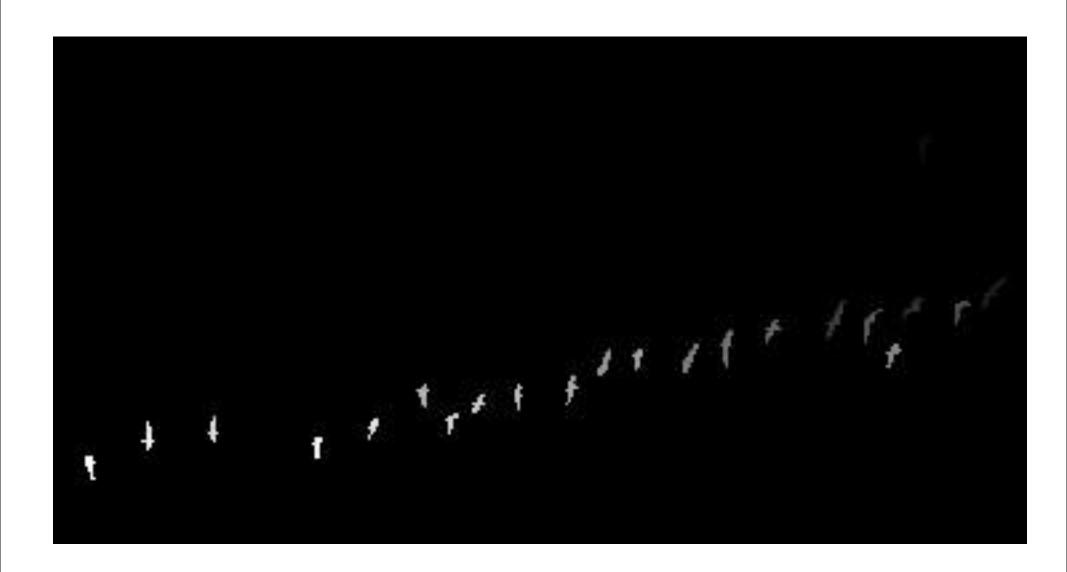
Goose Detector



Goose 4--components (26)

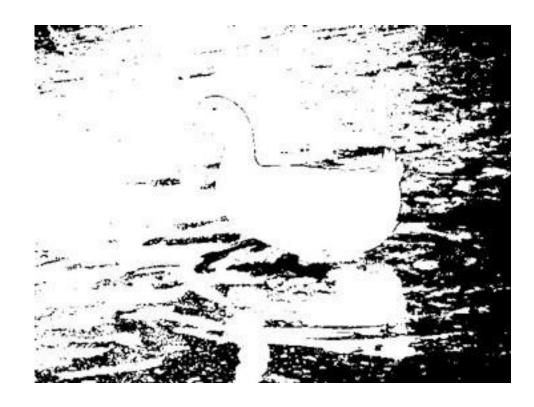


Goose 8--components (22)



Connected Components

What happens if we use the connected components algorithm on this image?



How might we improve our implementation?

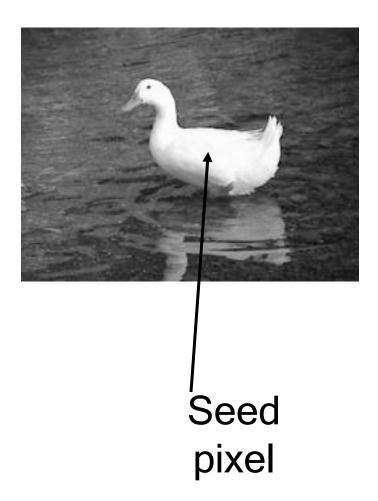
Region Growing

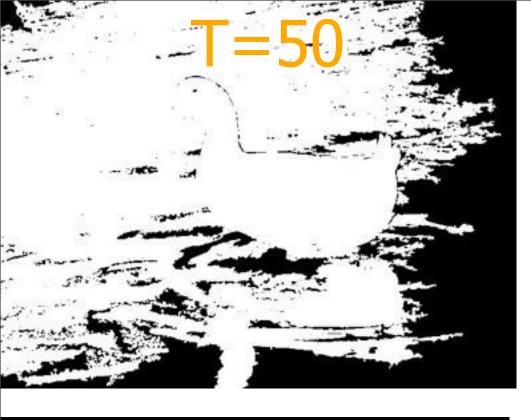
- Start from a seed point or region.
- Add neighbouring pixels that satisfy the criteria defining a region.
- Repeat until we can include no more pixels.

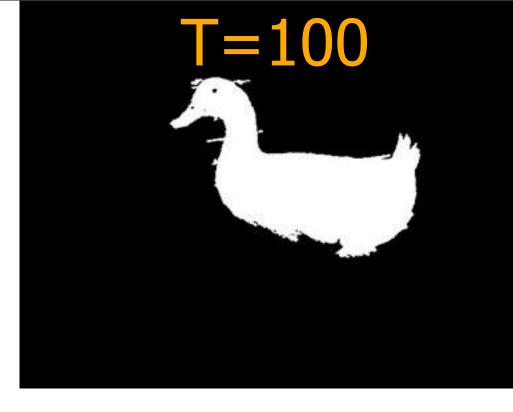
Region Growing example

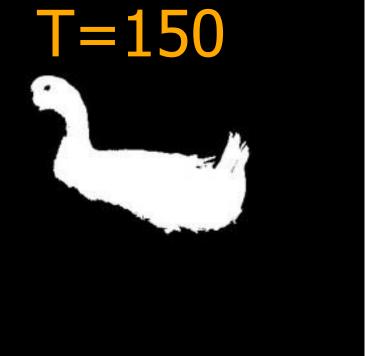
- Pick a single seed pixel.
- Inclusion test is a simple grey level threshold:

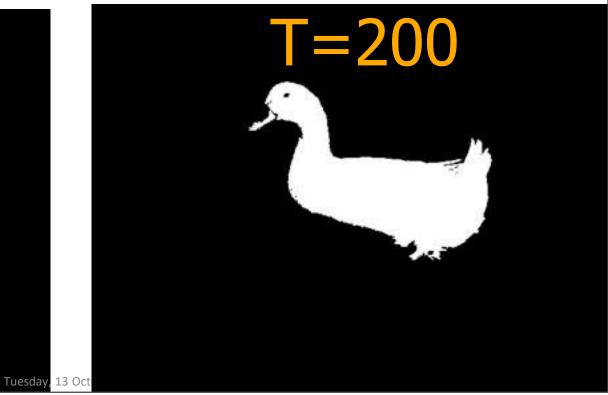
```
function test = include(p)
test = (p>=T);
```

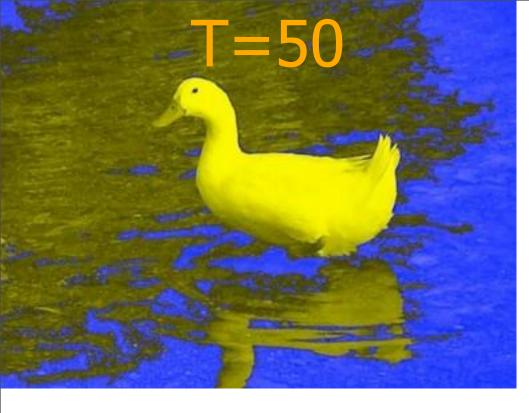


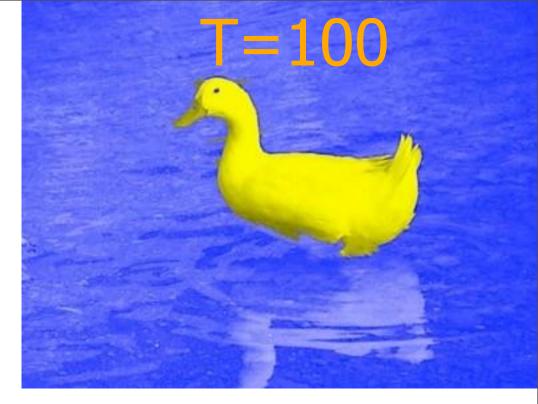


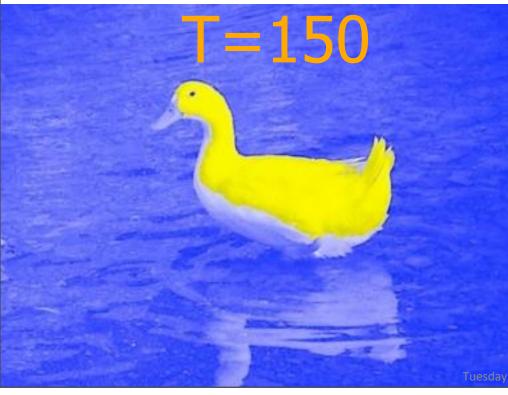


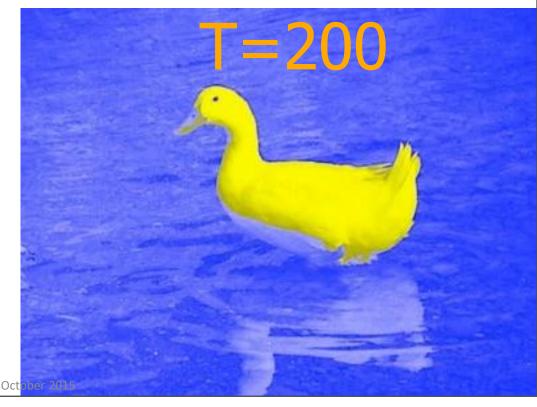






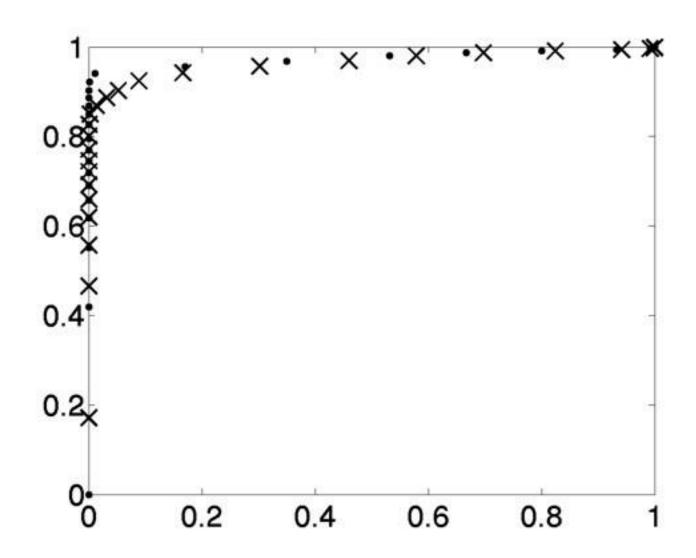






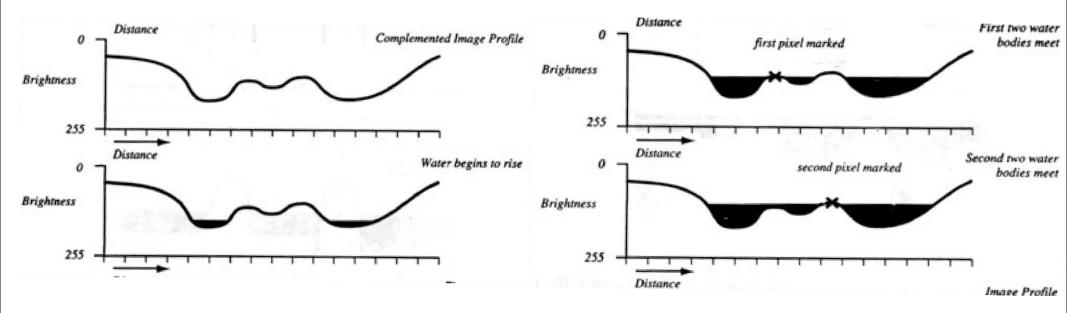
ROC Curve

- Region growing
- × Thresholding

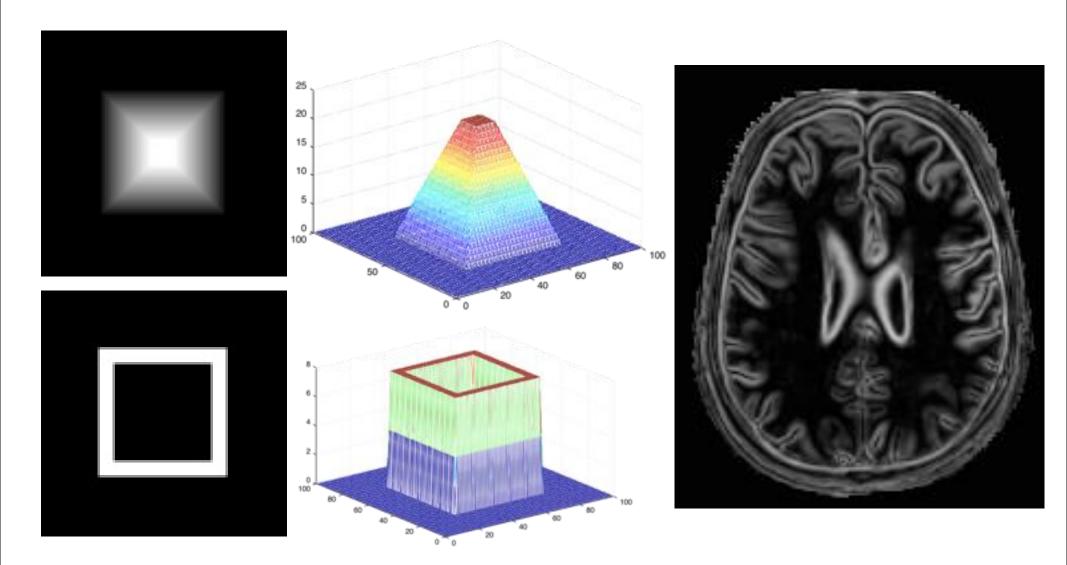


Watershed Algorithm

- The objective is to find watershed lines.
- Suppose that a hole is punched in each regional minimum and that the entire topography is flooded from below by letting water rise through the holes at a uniform rate.
- When rising water in distinct catchment basins is about to merge, a dam
 is built to prevent merging. These dam boundaries correspond to the
 watershed lines.



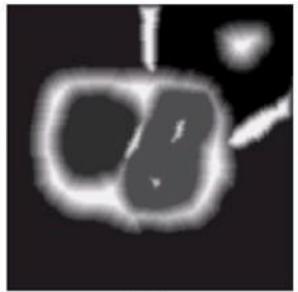
Watershed Segmentation



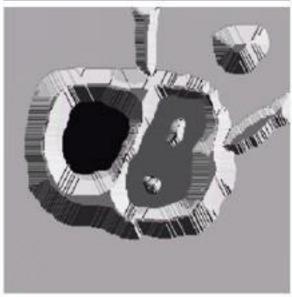
In this case, each object is distinguished from the background by its raised edges

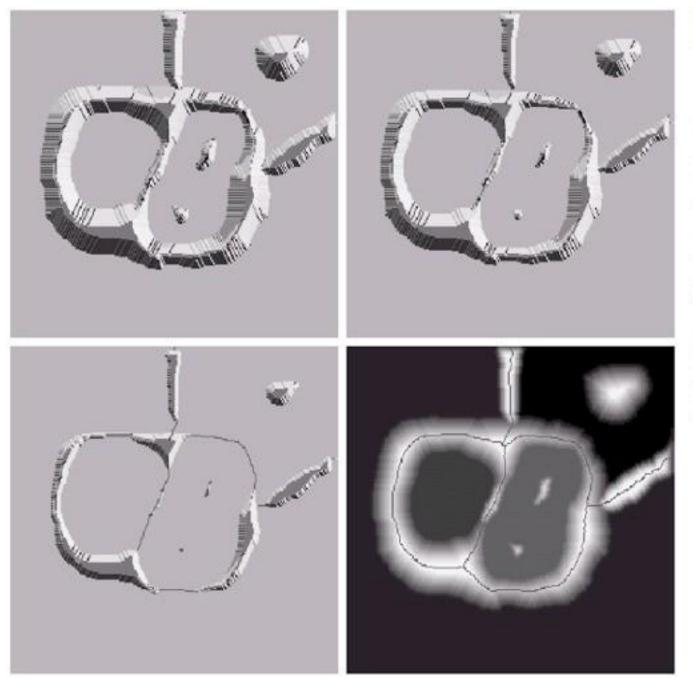
Basic Steps

- Piercing holes in each regional minimum of I
- The 3D topography is flooded from below gradually
- 3. When the rising water in distinct catchment basins is about to merge, a dam is built to prevent the merging









(Continued)
(e) Result of further flooding.
(f) Beginning of merging of water

from two

catchment basins (a short dam was built between

them). (g) Longer dams. (h) Final watershed

(segmentation) lines. (Courtesy of Dr. S. Beucher, CMM/Ecole des Mines de Paris.)

4. The dam boundaries correspond to the watershed lines to be extracted by a watershed segmentation algorithm

Watershed Algorithm

```
Set label at every pixel to -1

For each seed point u

For each neighbour v of u

Label[v] = label [u]

Add v to priority queue: priority by gray level

while(elements in queue)

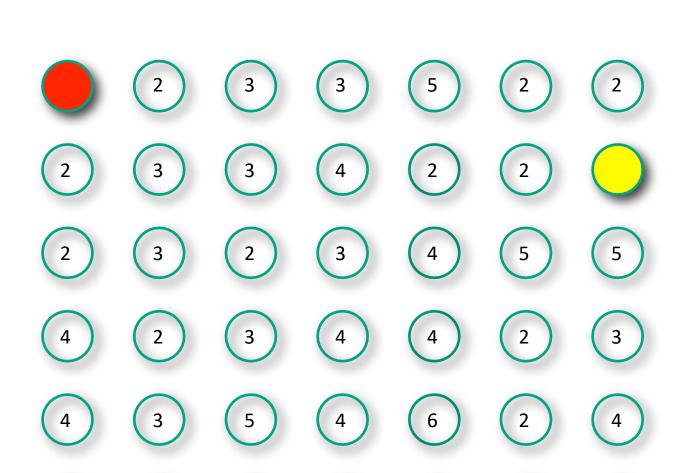
u = popMinimum(Queue);

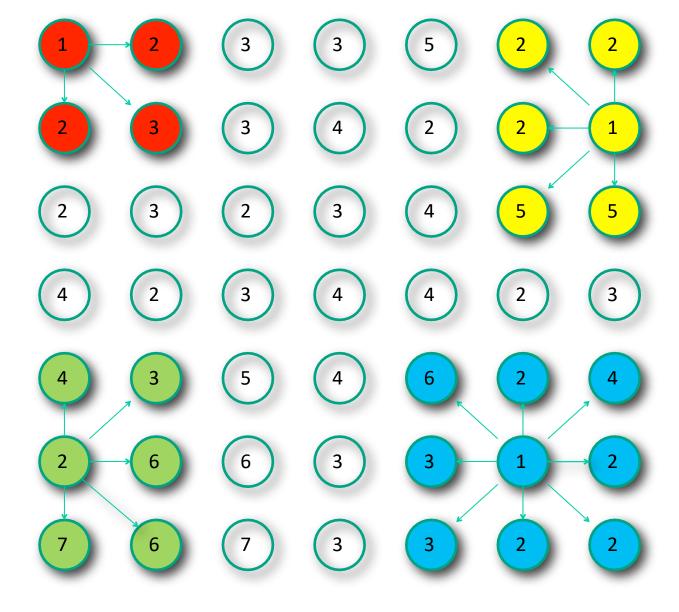
For each neighbour v of u

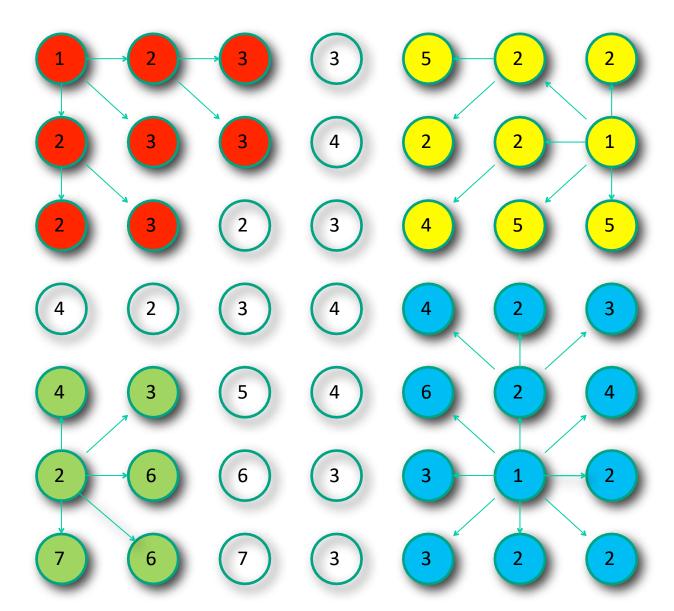
If (label[v]!==-1)

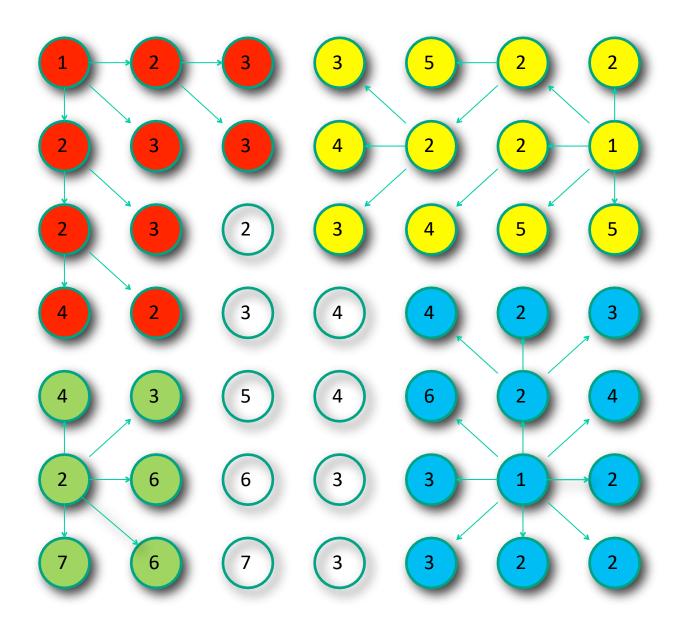
Label[v] = label [u]

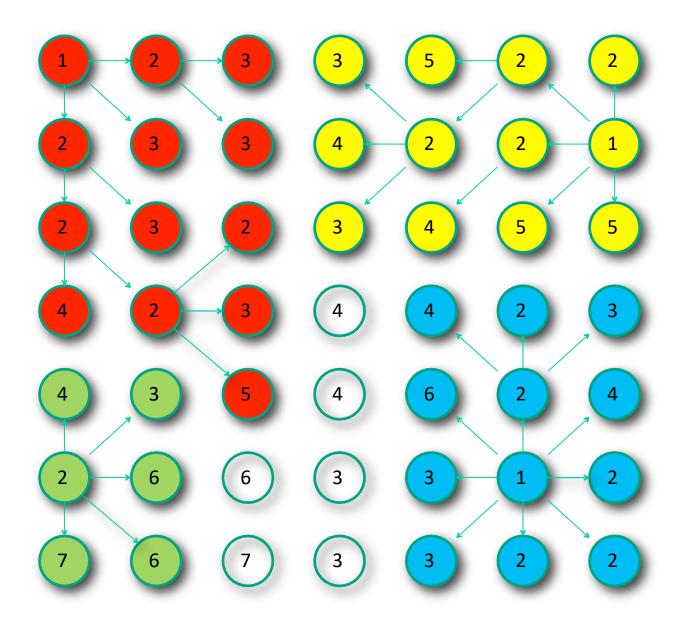
Add v to priority queue: priority by gray level
```

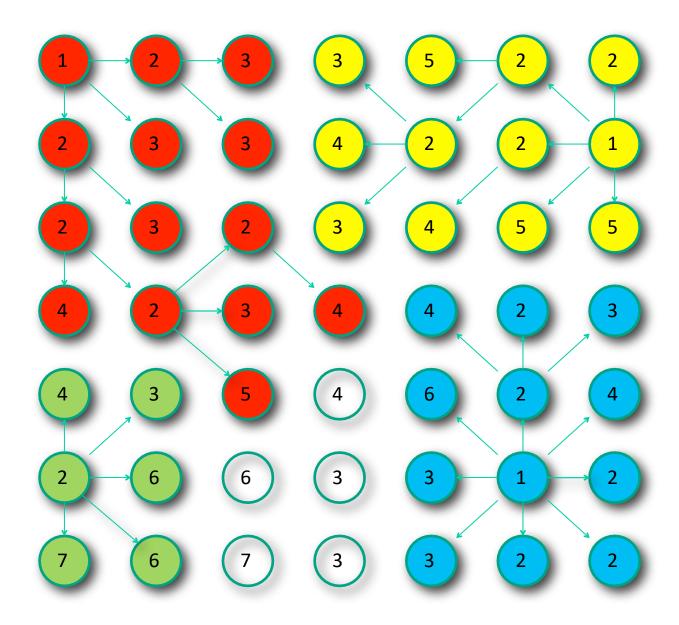


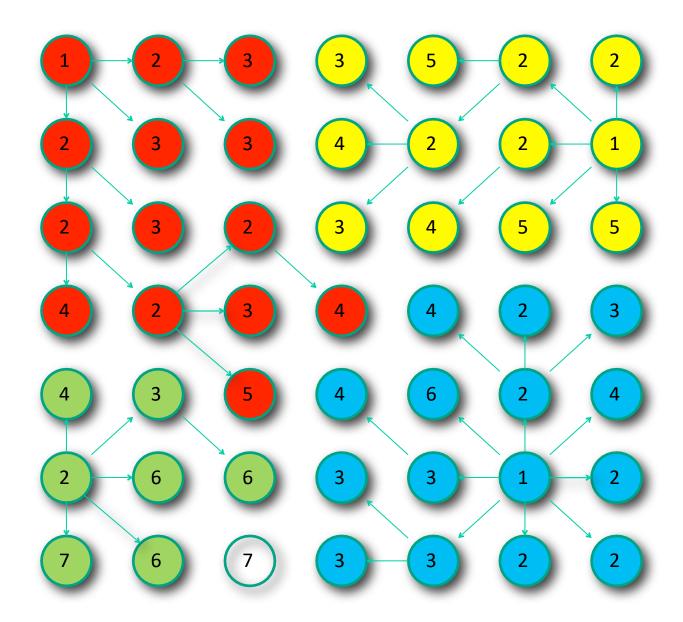


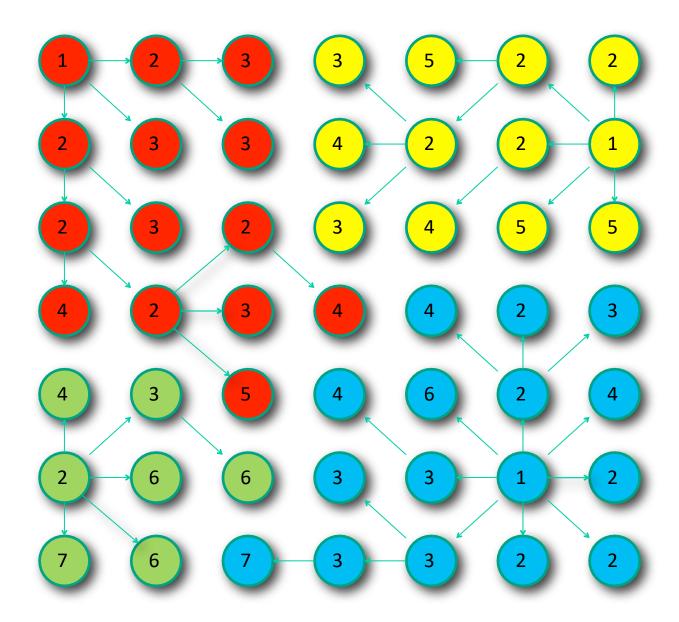


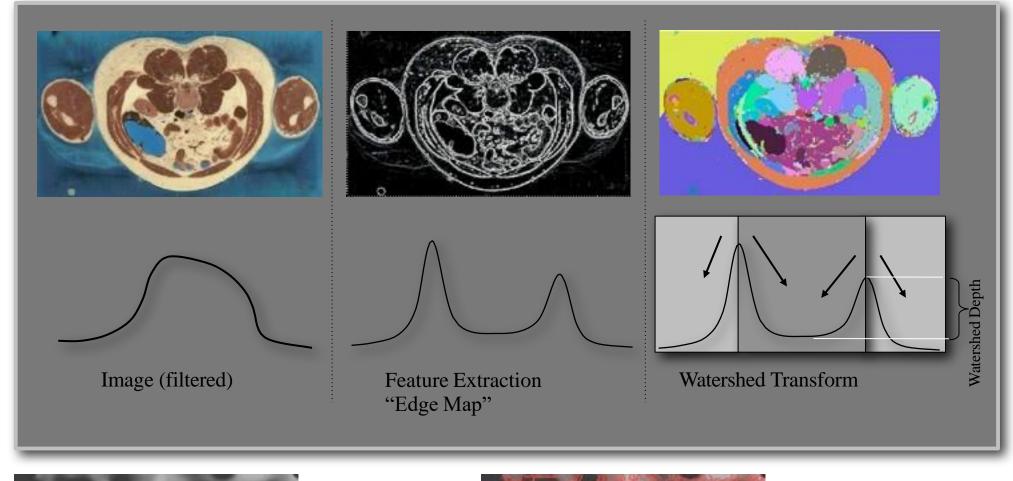


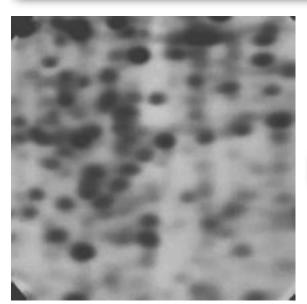


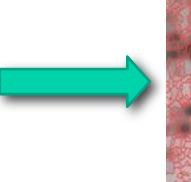


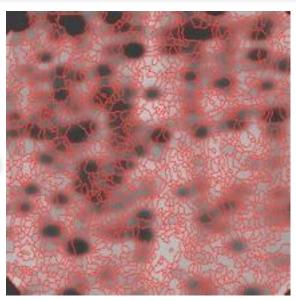






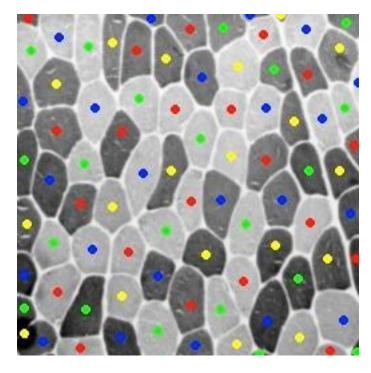






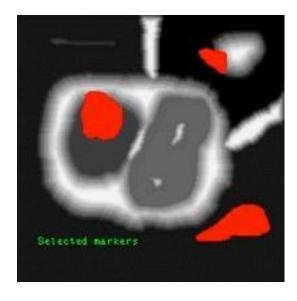
Problem:
Watershed
algorithm
typically oversegments

Initialization



IDEA: Initialize regions using other information (cell nuclei) or by hand--marking.

Pre-defines number of segments









Analysis (pros & cons)

Thresholding

- Relies on a single global threshold
- Produces separate regions (may prefer connected components)

Region growing

- Supervised, requires 1+ seed point
- Relies on a single threshold
- Produces one region

Watershed

- May be supervised or unsupervised
- No need for threshold parameter
- Partitions image into many regions
- To get good results best to specify number and positions of seeds

What is image segmentation?

- 1. Thresholding, region labelling and growing algorithms
 - (connected components, region growing, watershed)
- 2. Statistical Segmentation
 - (k-means, mean shift)
- 3. Graph based methods
 - (Merging algorithms, split/merge)
- 4. Edge based methods
 - (Intelligent Scissors, Snakes)

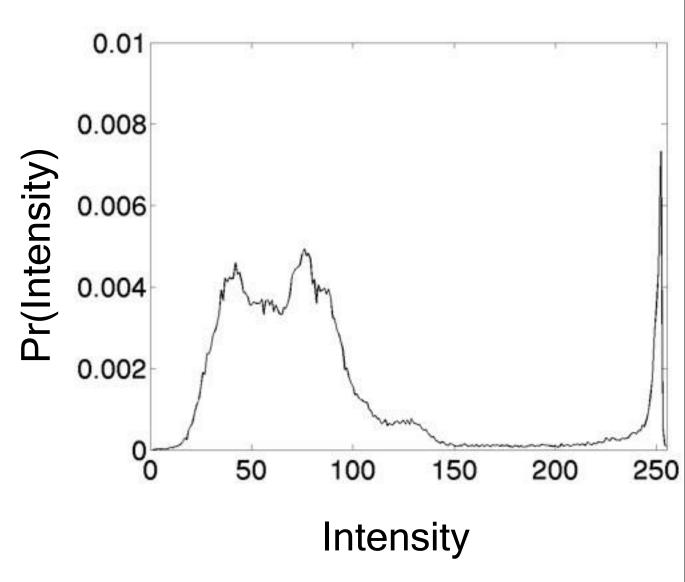
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Choosing Threshold Automatically



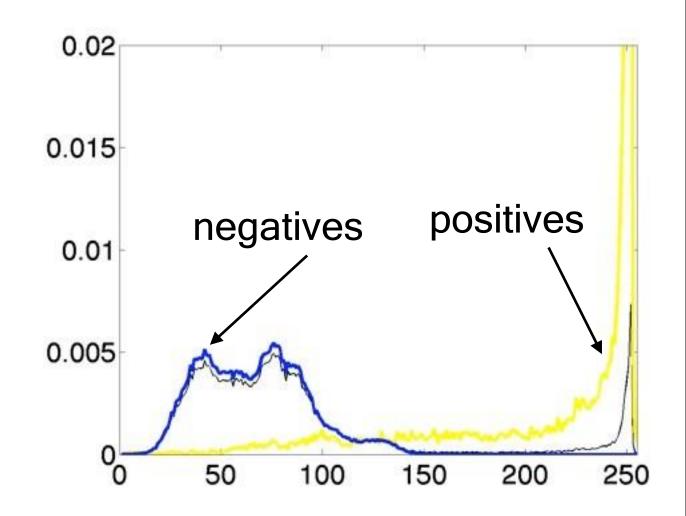
Q. For binary thresholding, how can we choose the threshold automatically?



Choosing Threshold Automatically

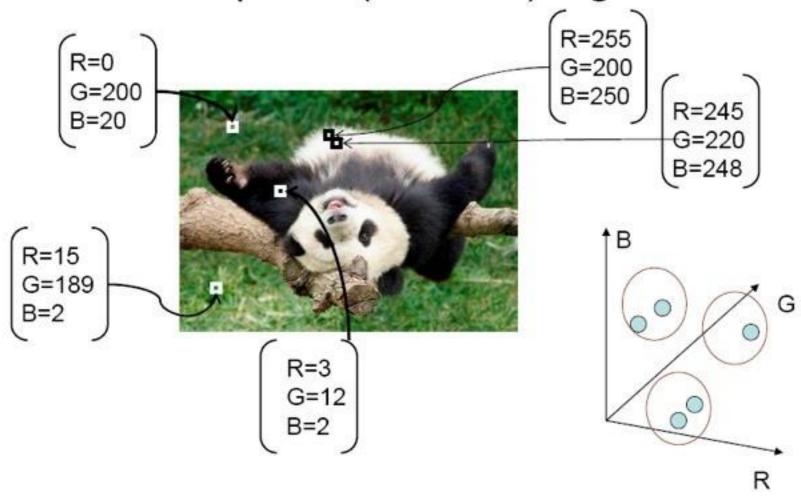


Q. For binary thresholding, how can we choose the threshold automatically?



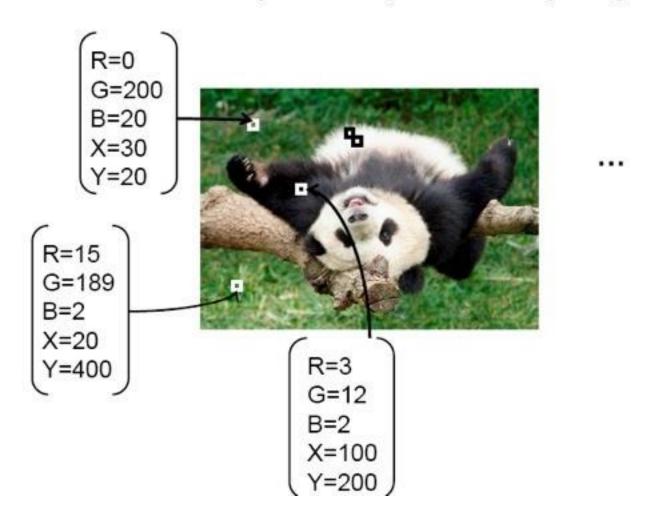
Segmentation as clustering

Cluster similar pixels (features) together



Segmentation as clustering

Cluster similar pixels (features) together



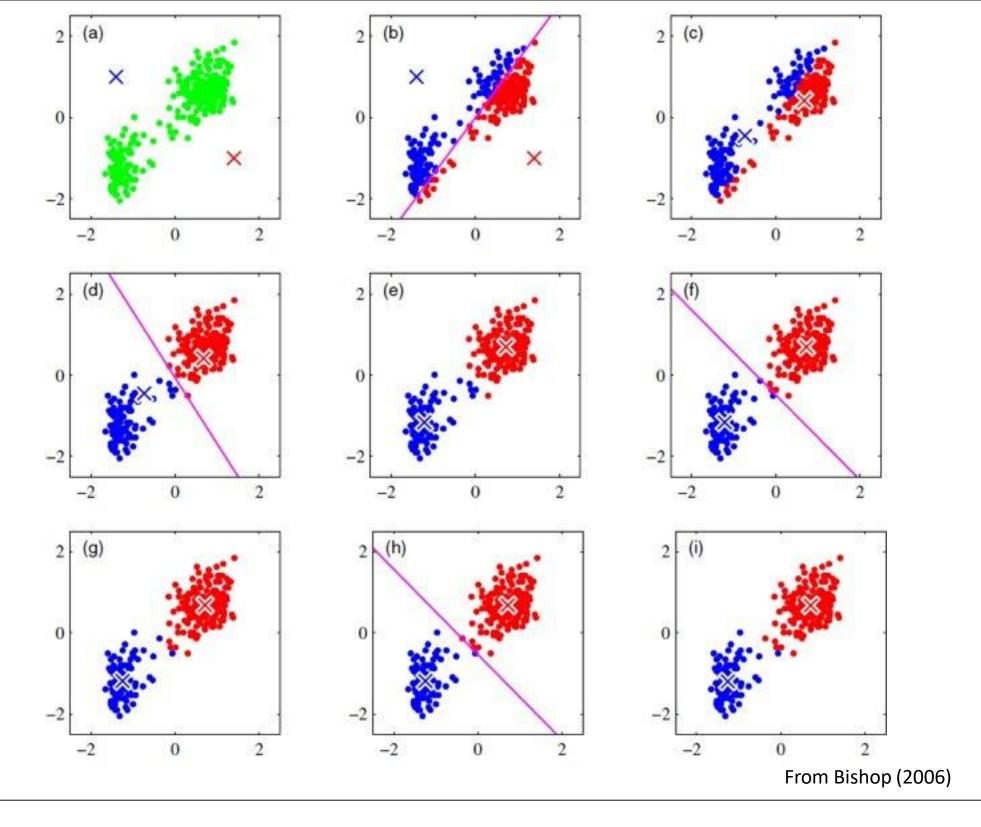




Figure 9.3 Two examples of the application of the K-means clustering algorithm to image segmentation showing the initial images together with their K-means segmentations obtained using various values of K. This also illustrates of the use of vector quantization for data compression, in which smaller values of K give higher compression at the expense of poorer image quality.

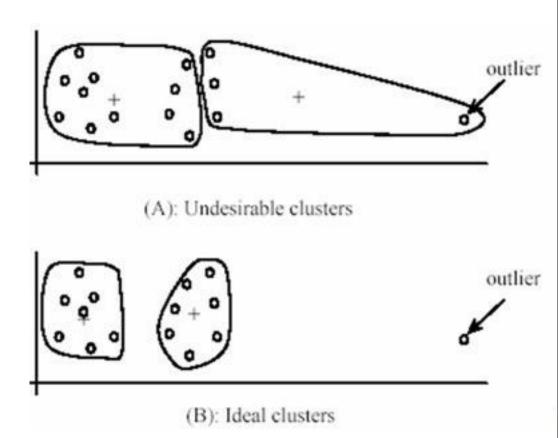
K-Means

Pros

- Simple and fast
- Converges to a local minimum of the distance function

Cons

- Need to pick K
- Sensitive to initialization
- Sensitive to outliers
- Only finds "spherical" clusters



Comparison

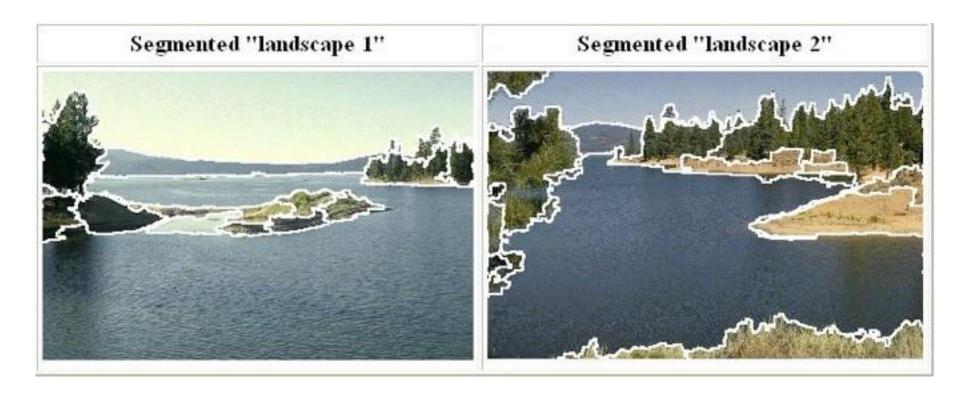
Watershed

- May be supervised or unsupervised
- No need for threshold parameter
- Partitions image into many regions
- To get good results best to specify number and positions of seeds

K-Means

- Unsupervised
- Works in high--dimensions
- May produce disjoint segments
- Must specify number of clusters but not position

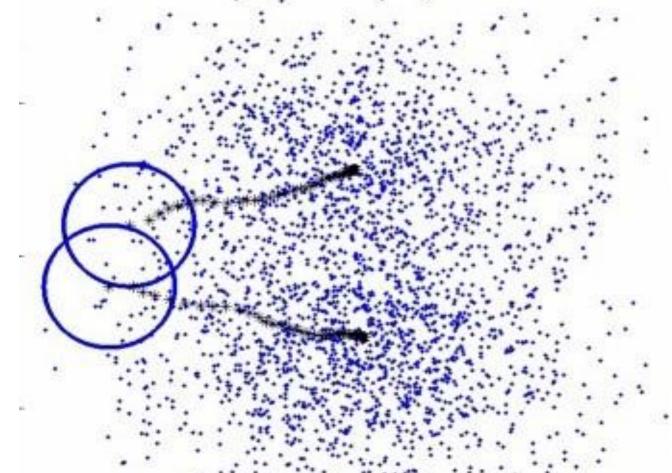
Mean shift Segmentation

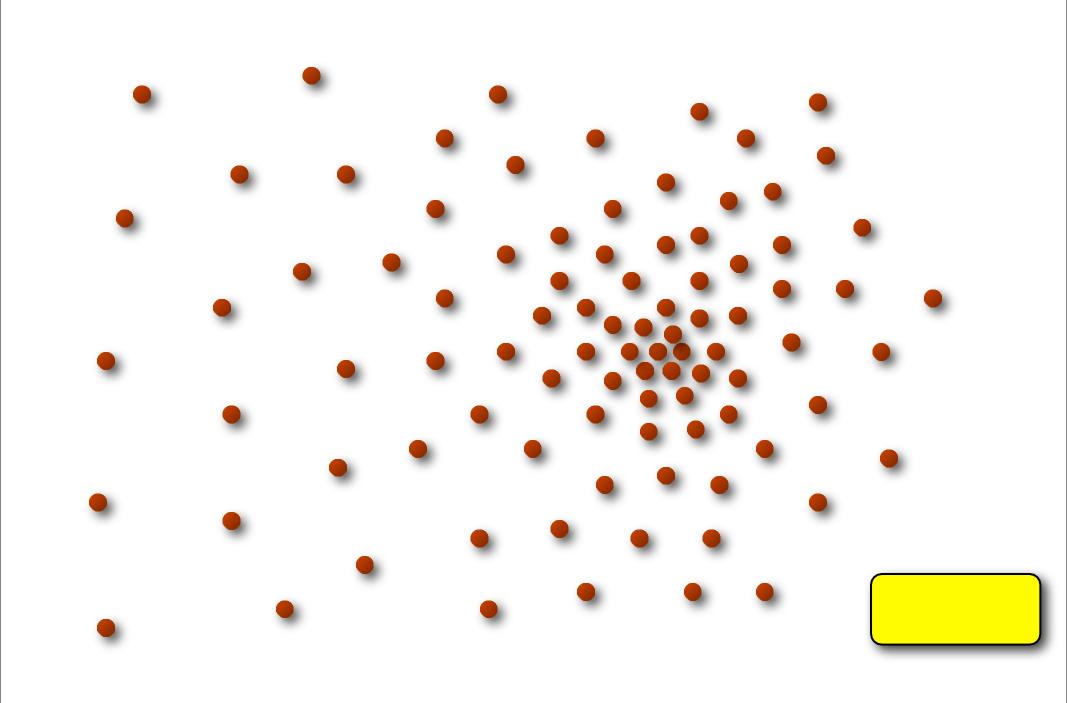


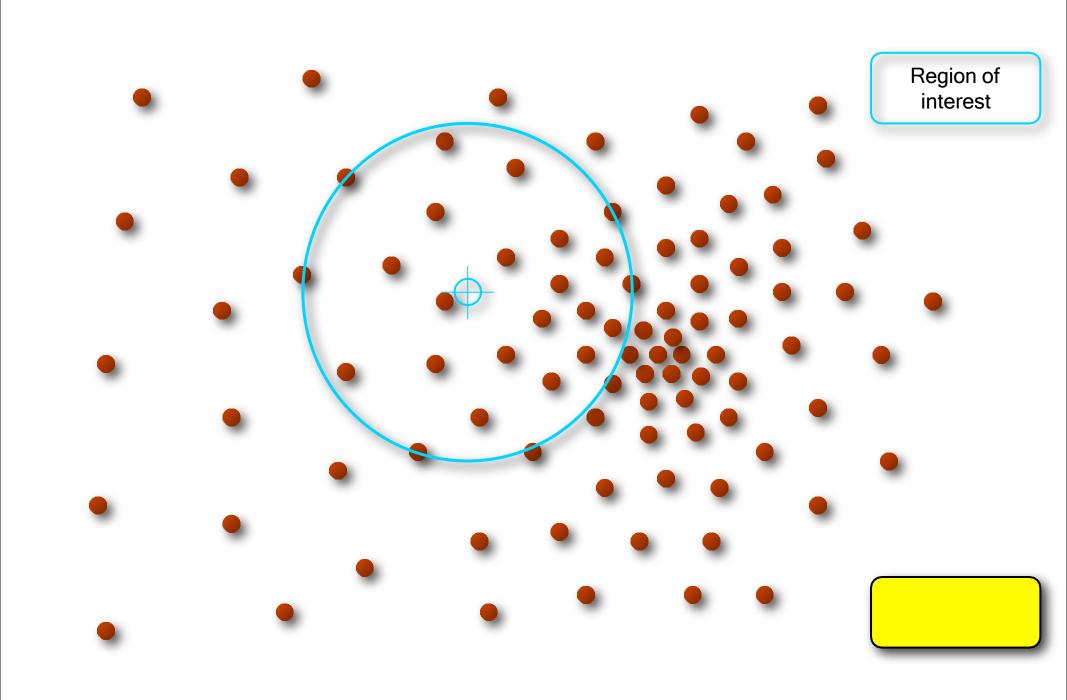
Mean Shift: A Robust Approach toward Feature Space Analysis, PAMI 2002.

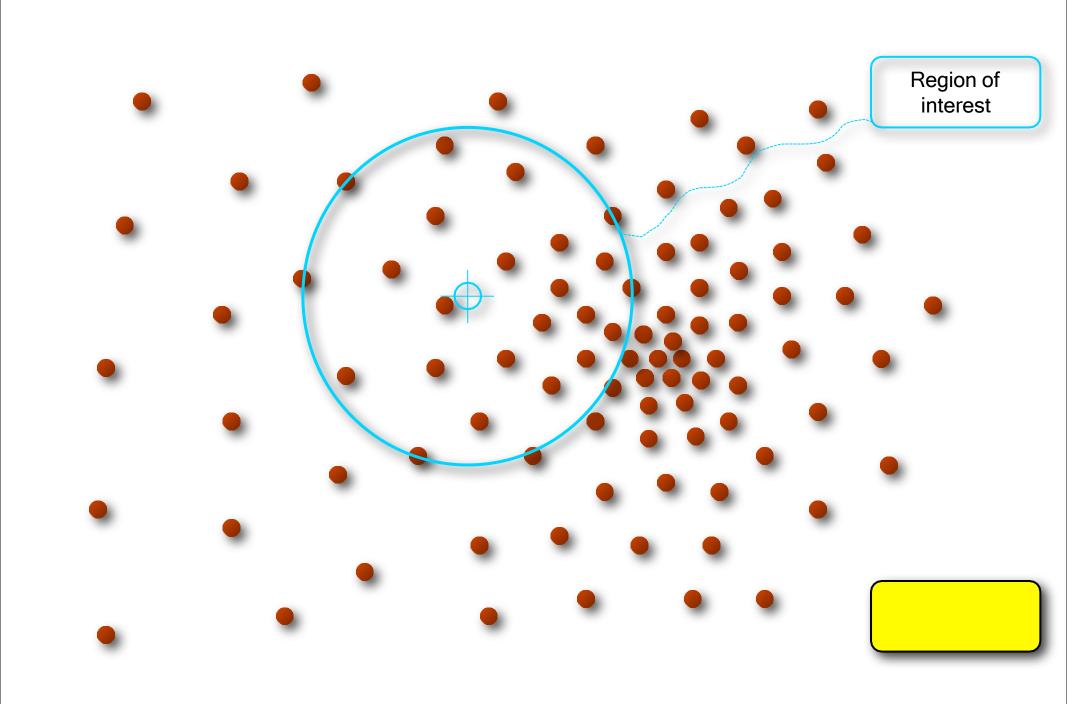
D. Comaniciu and P. Meer

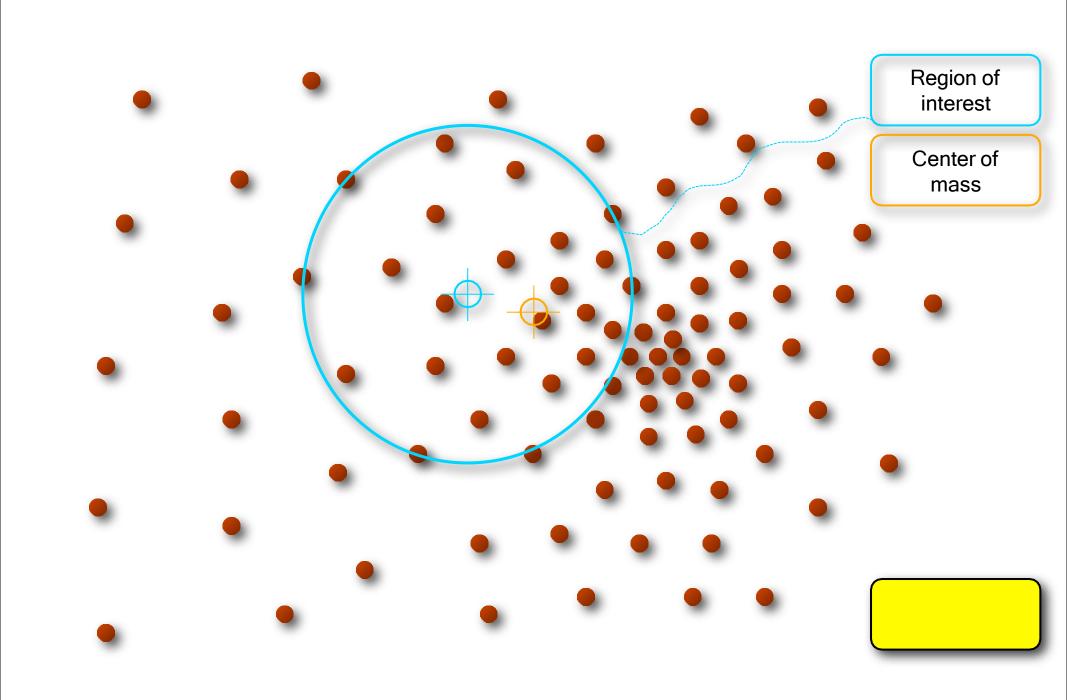
- The mean shift algorithm seeks a *mode* or local maximum of density of a given distribution
- Choose a search window (width and location)
- Compute the mean of the data in the search window
- Center the search window at the new mean location
- Repeat until convergence

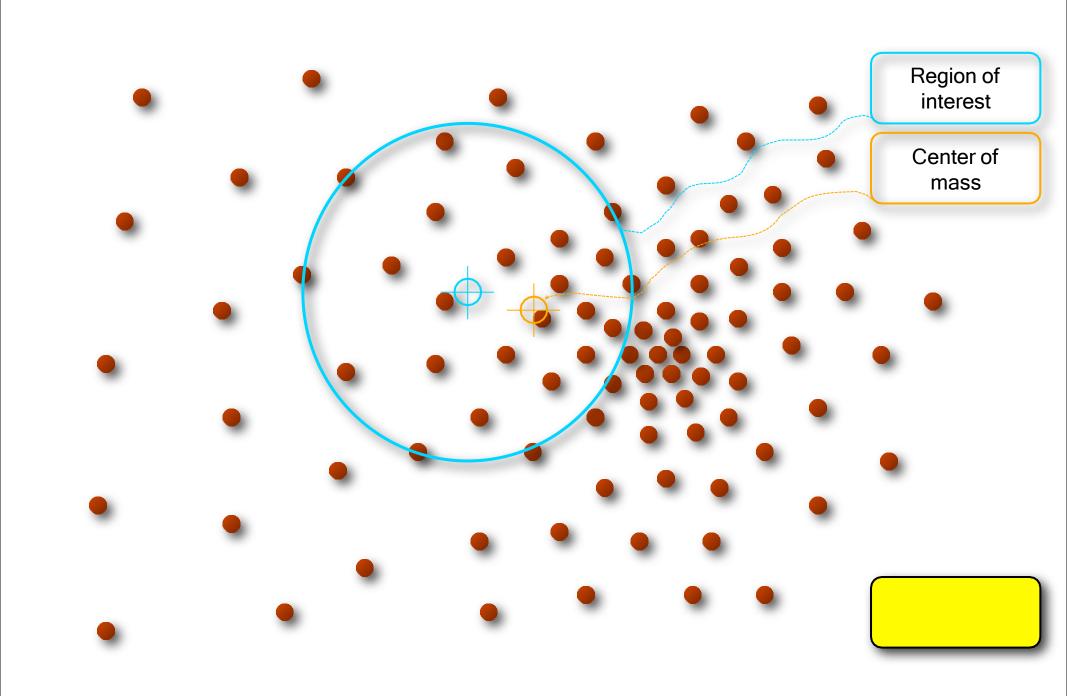


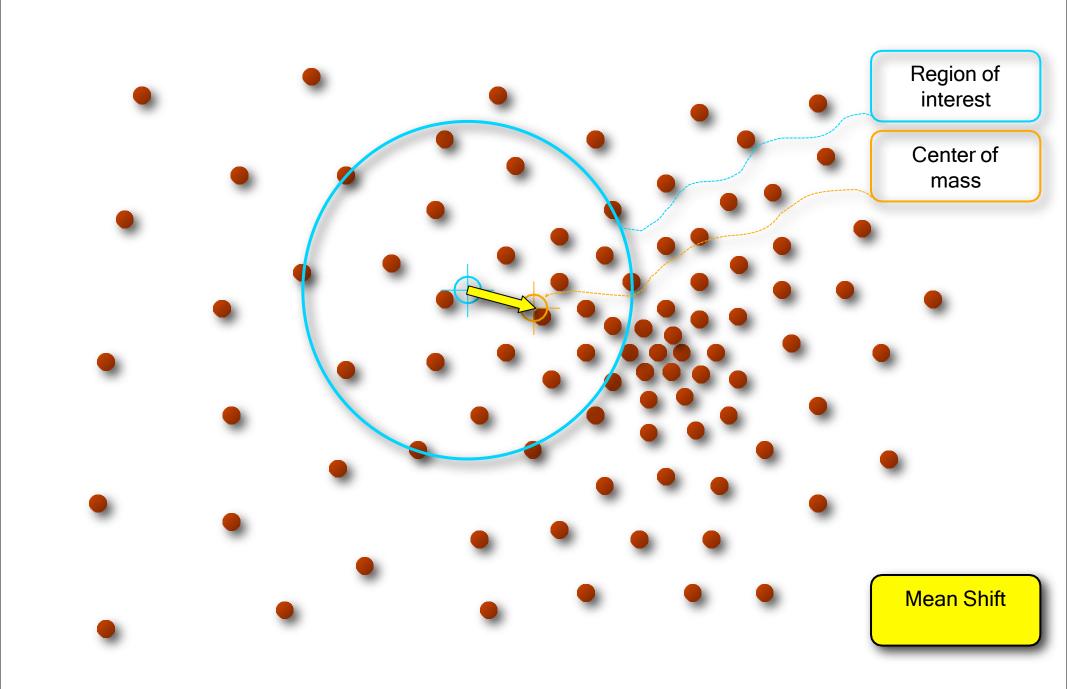


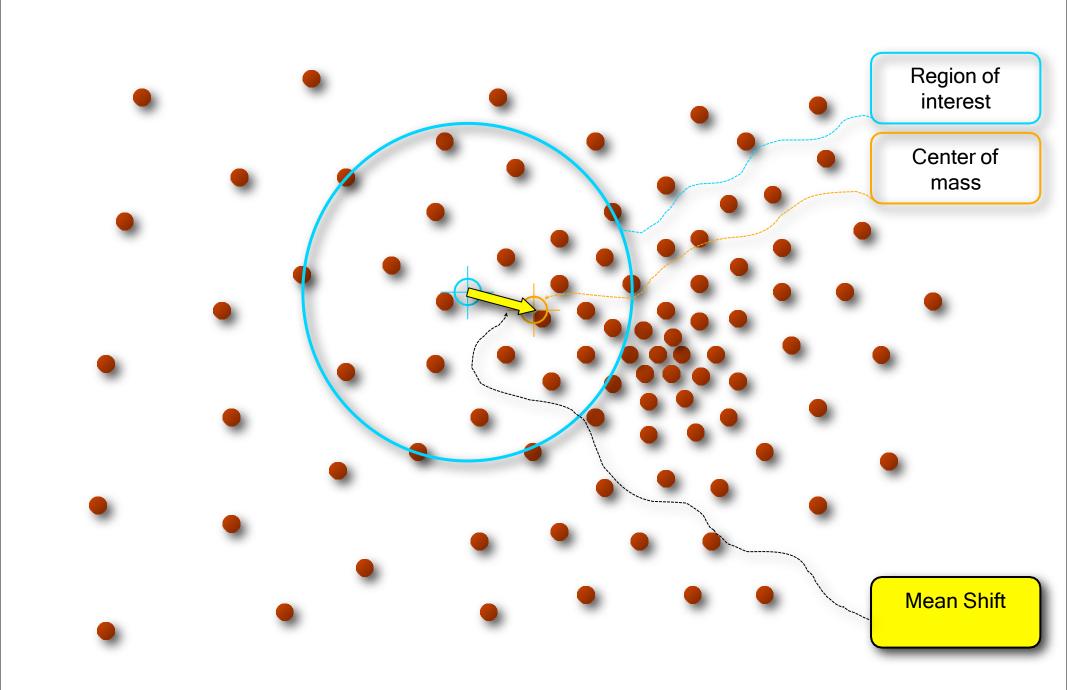


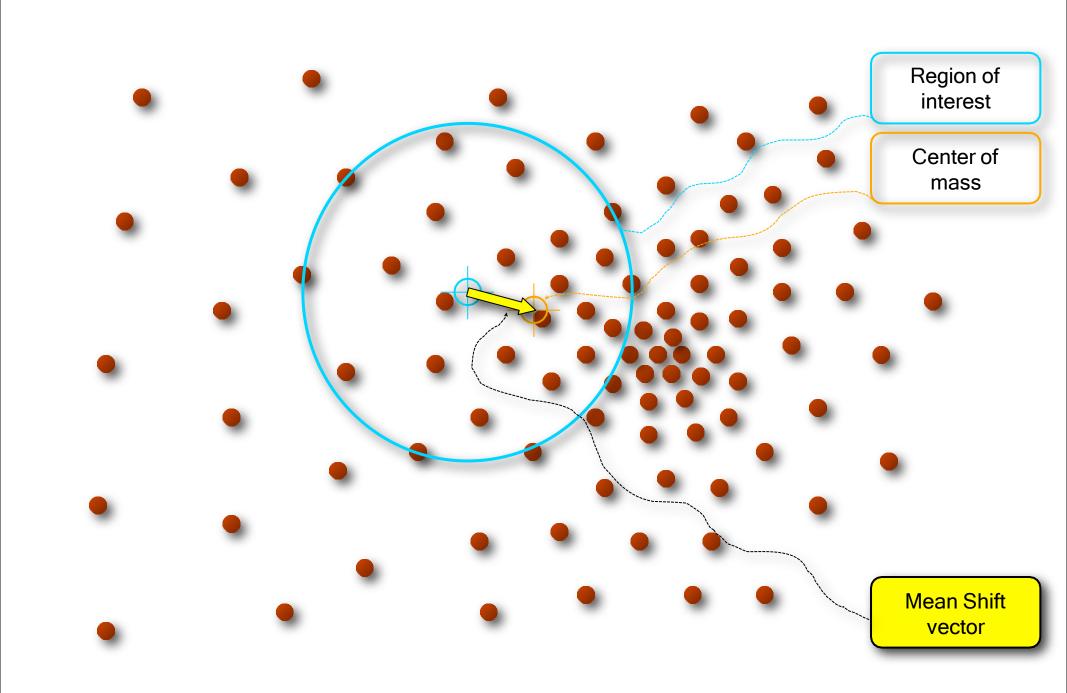


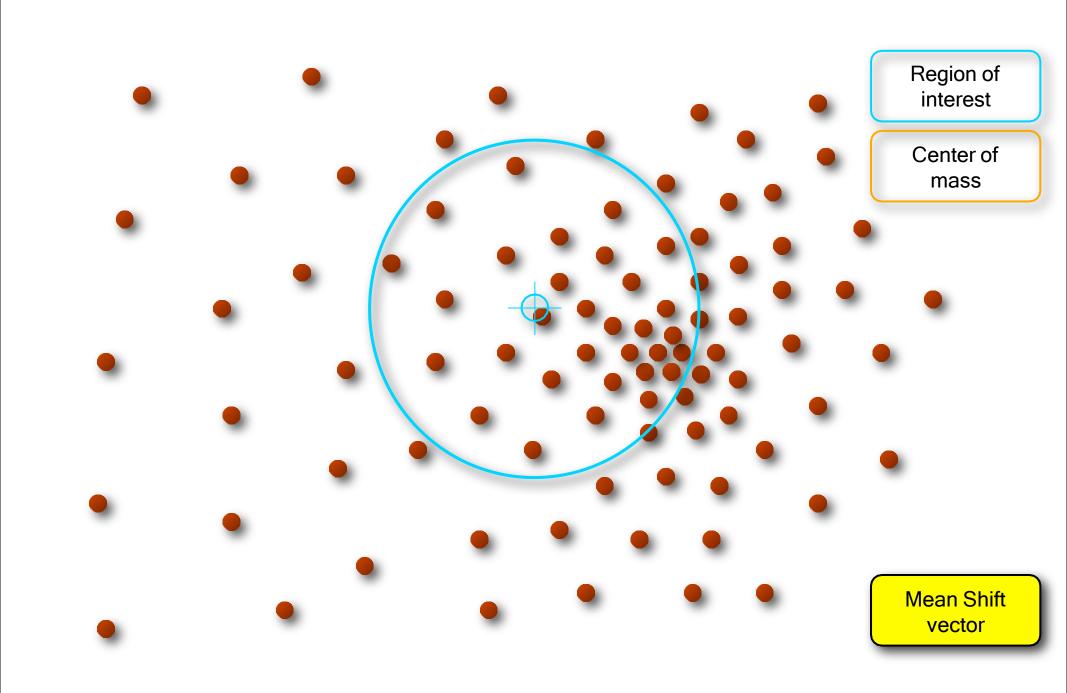


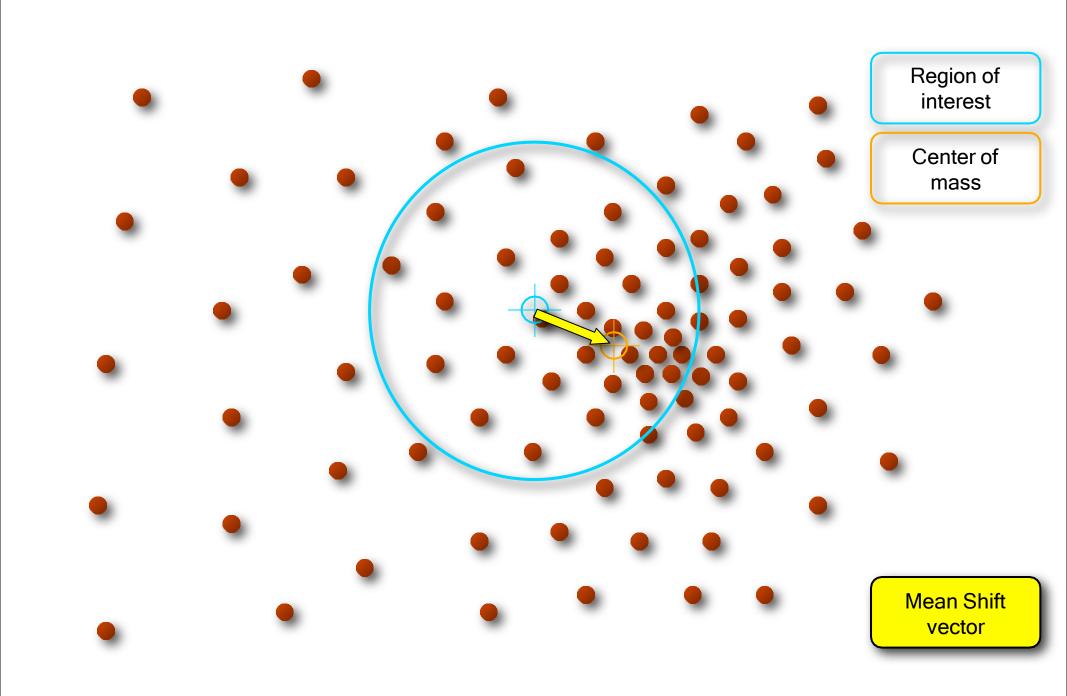


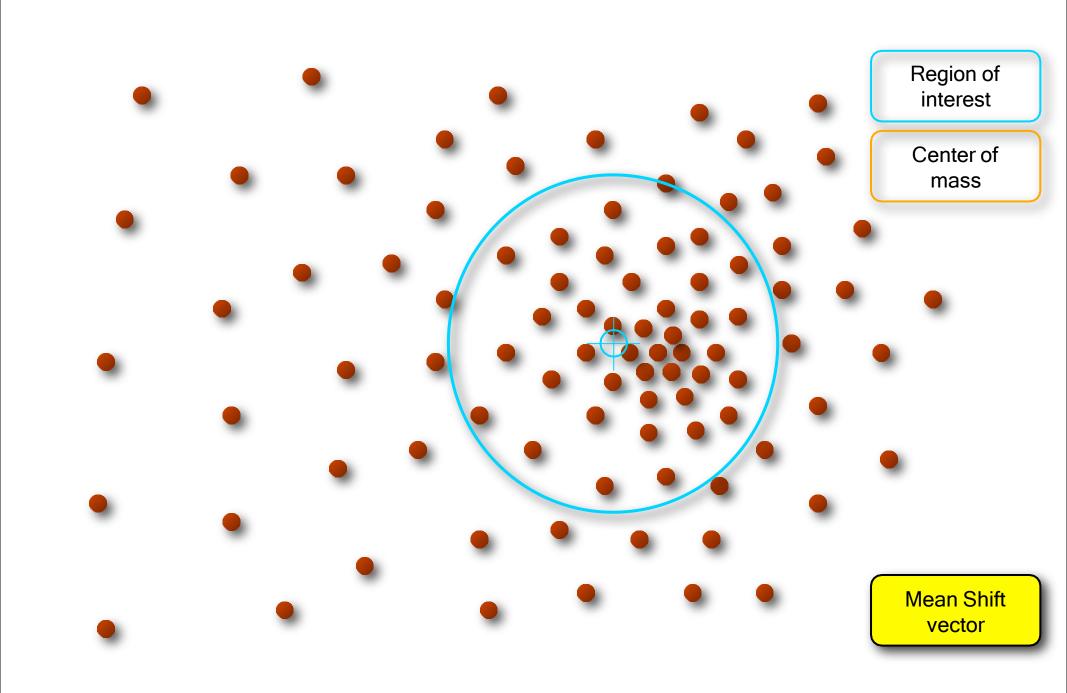


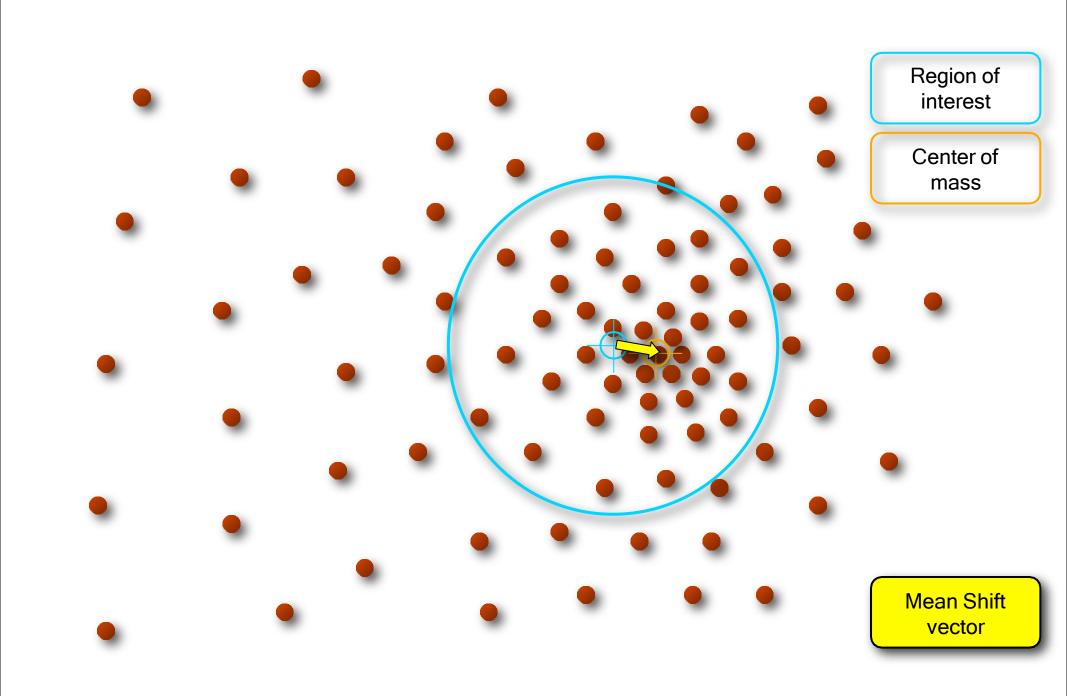


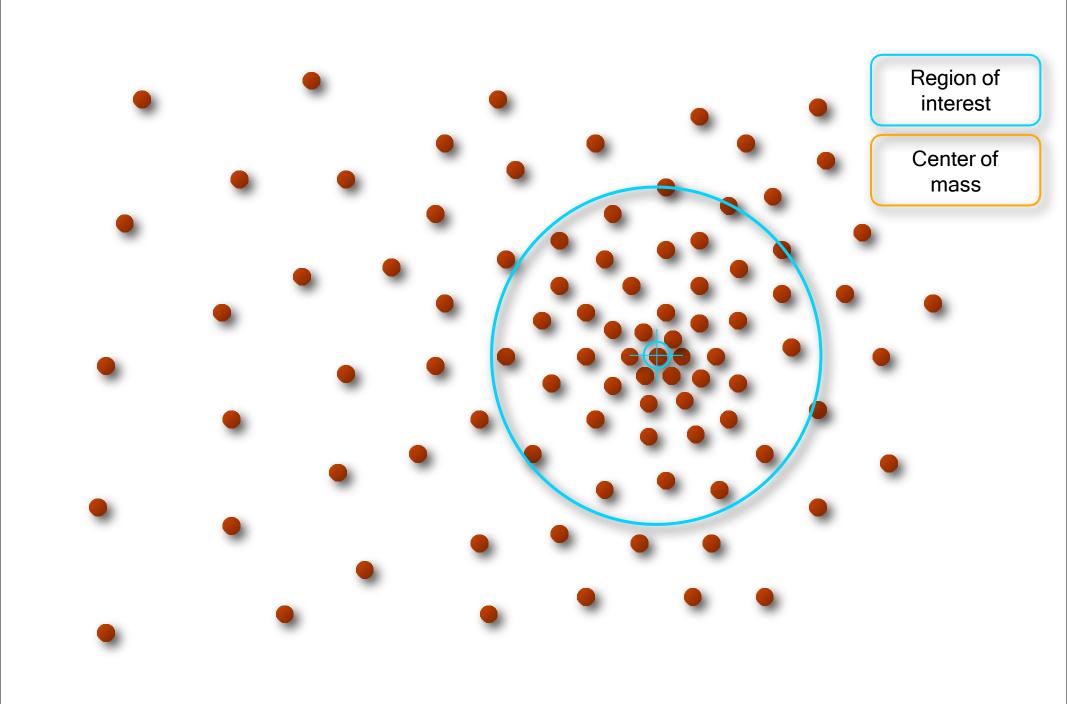


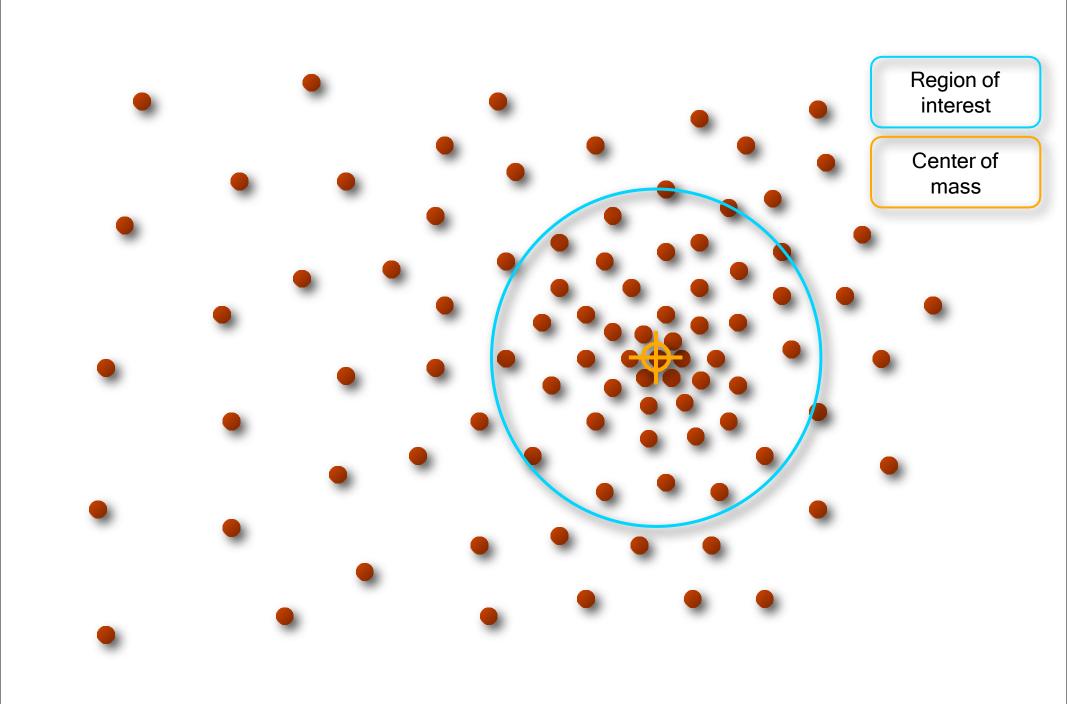






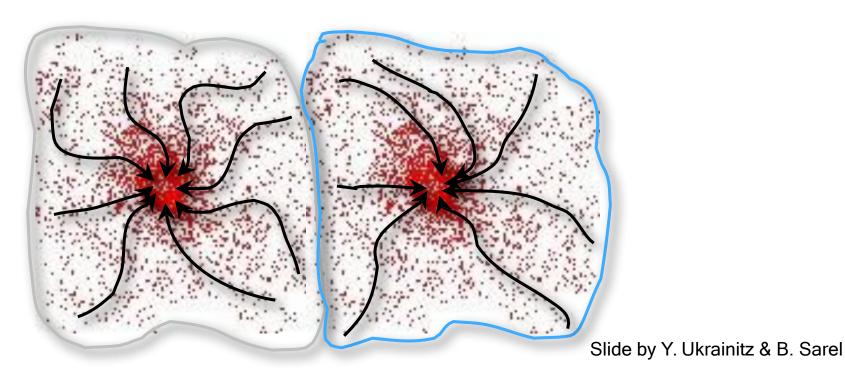






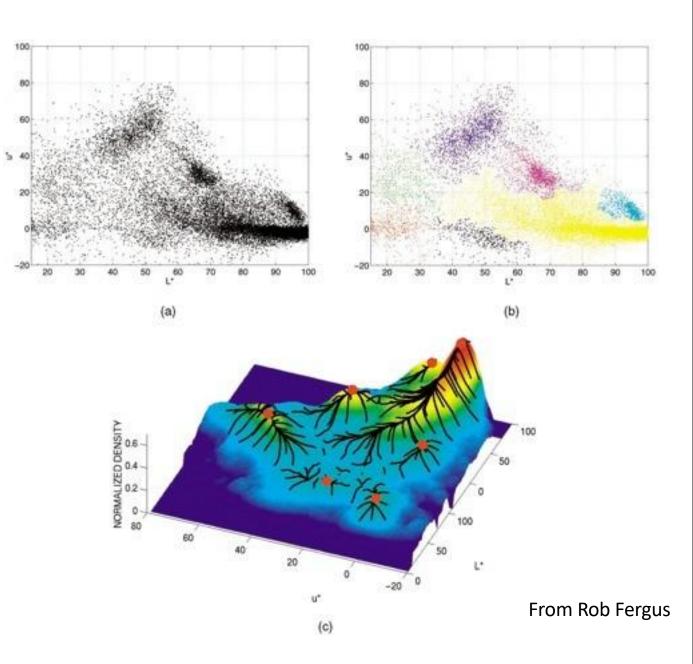
Mean shift Clustering

- 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

- 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



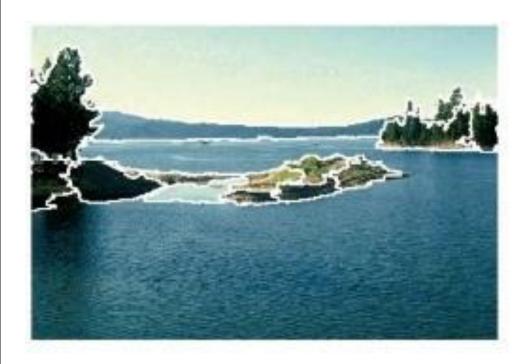
Mean-shift Clustering Result

















More Results



Mean shift Clustering

Pros

- Does not assume spherical clusters
- Just a single parameter (window size)
- Finds variable number of modes
- Robust to outliers

Cons

- Output depends on window size
- Computationally expensive
- Does not scale well with dimension of feature space

Comparison

Watershed

- May be supervised or unsupervised
- No need for threshold parameter
- Partitions image into many regions
- To get good results best to specify number and positions of seeds

K-Means

- Unsupervised
- Works in high--dimensions
- May produce disjoint segments
- Must specify number of clusters but not position

Mean-Shift

- Unsupervised
- Works in high--dimensions (kind of)
- May produce disjoint segments
- "Only" specify windows size

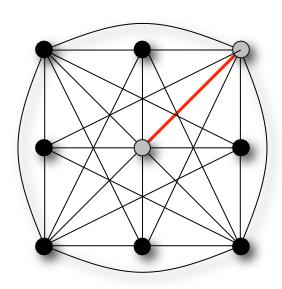
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 - (Intelligent Scissors, Snakes)

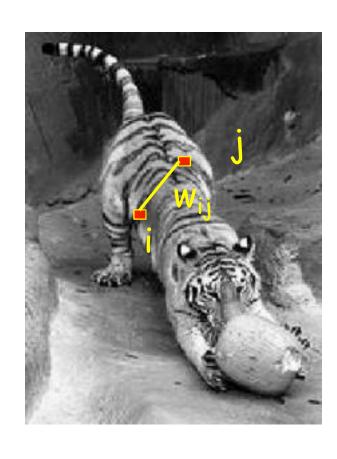
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 - (Intelligent Scissors, Snakes)

Images as Graphs



Similar pixels

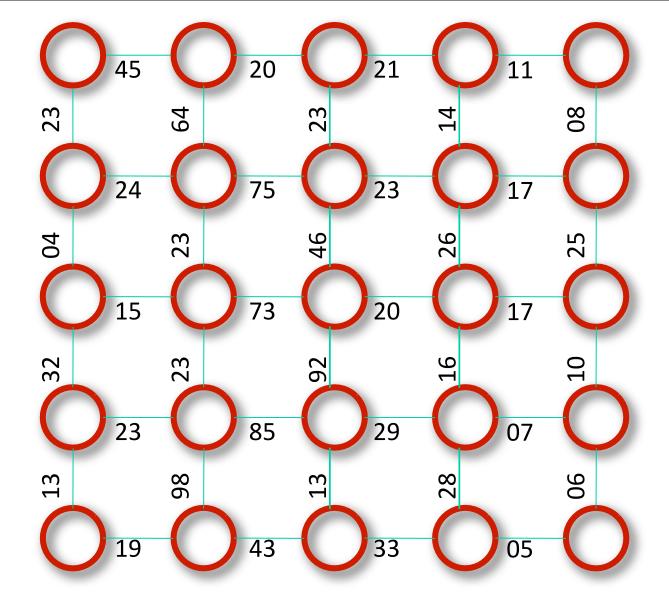


- Node for every pixel
- Edge between every pair of pixels (or every pair of "sufficiently close" pixels)
- Each edge is weighted by the affinity or similarity of the two nodes

Source: S. Seitz

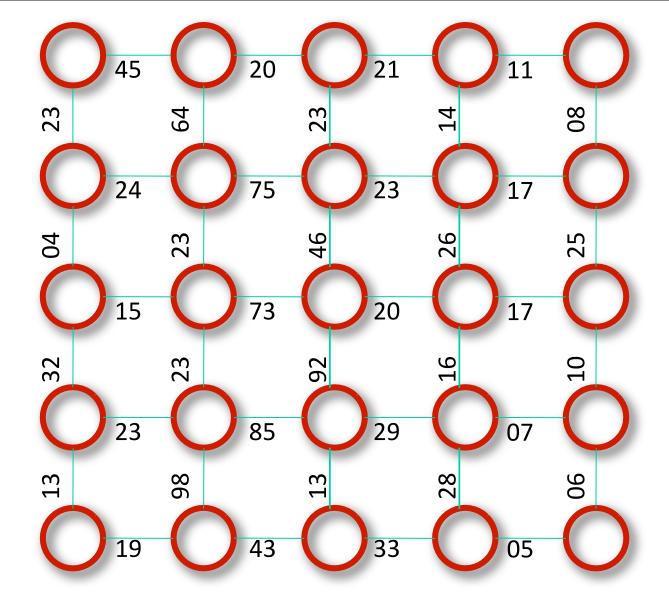
Greedy Merging Algorithm

- 1. Initialize each pixel to a separate segment
- 2.Assign a cost, c_{ij} to each edge (possibly based on pixel difference)
- 3. Sort all edges by edge strength
- 4.Connect edges in order of strength, merging segments as we go
- 5.Stop when we have reached some threshold edge strength

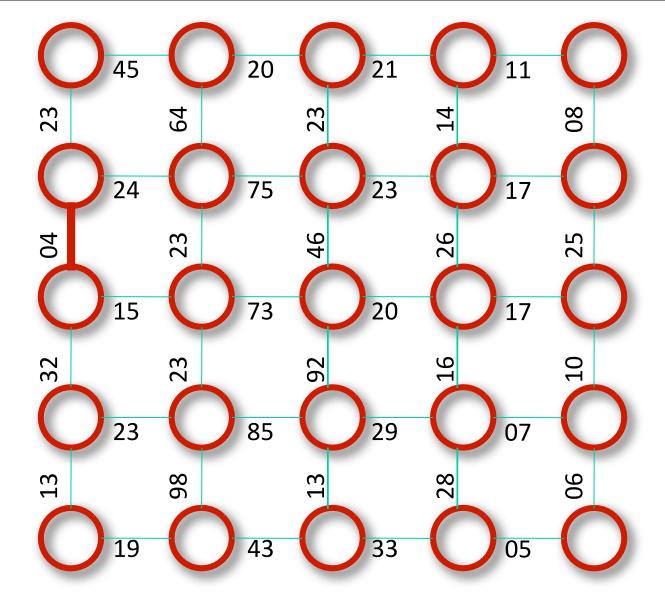


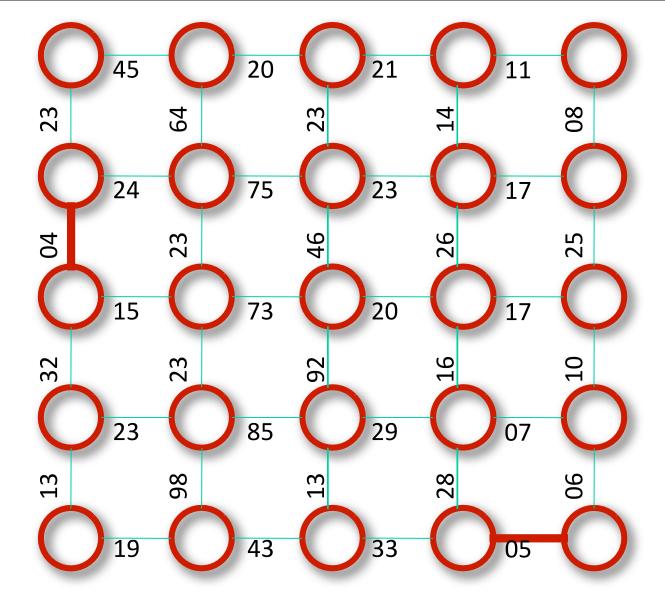
Extract all edge strengths

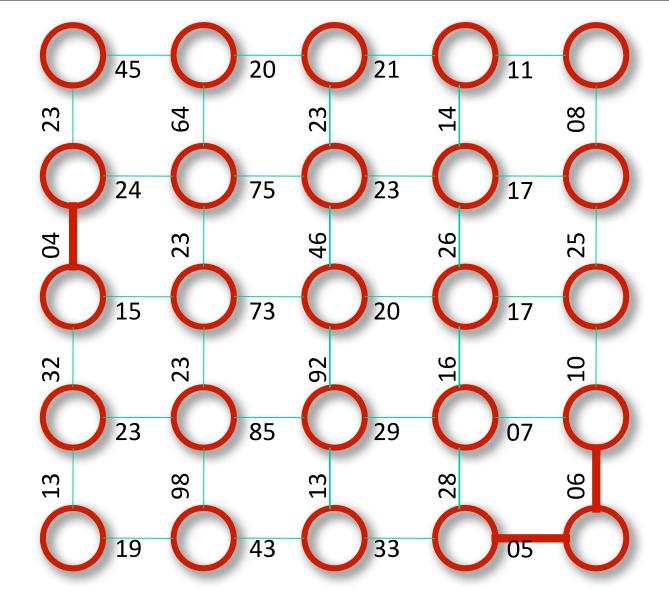
45, 20, 21, 11, 24, 75, 23, 17, 15, 73, 20, 17, 23, 85, 29, 07, 19,43,33, 05, 13, 22, 04,23, 98, 23, 23, 64, 13, 92, 46, 23, 28, 16, 26, 14, 06, 10, 25, 08

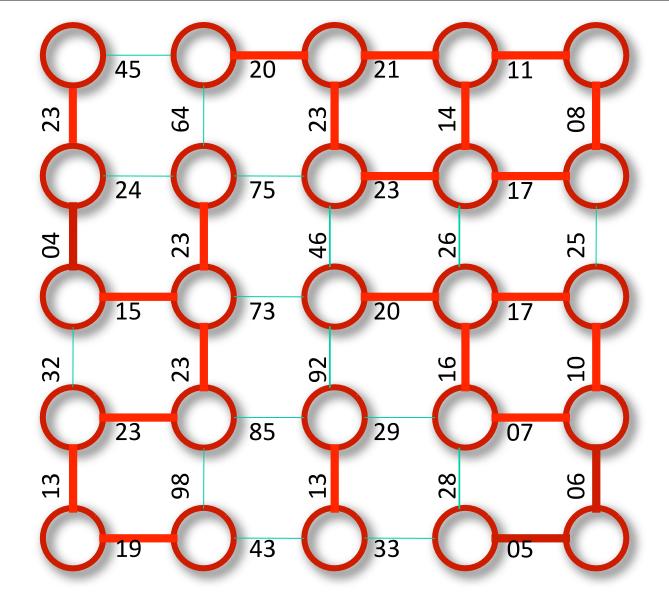


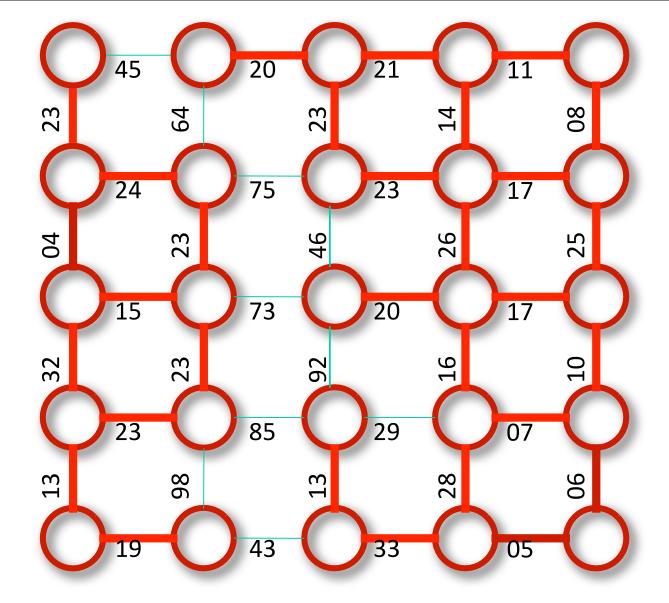
Sort edge strengths (remembering where they came from)











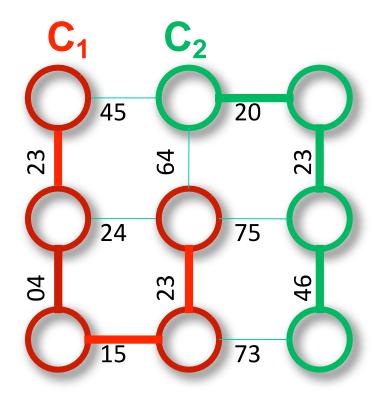
A Better Merging Algorithm

Efficient Graph--Based Image Segmentation (Felzenswalb, Huttenlocher [IJCV 2004])

- Assign weight w(e_{ij}) to each edge e_{ij}
- Sort edges by weight and run through
- Define Int(C) to be the maximum cost weight in the minimum spanning tree of cluster C
- Define Dif(C1,C2) to be lowest cost link joining components
- Merge components if

Dif
$$(C_{1,}C_{2}) < \min(\text{Int}(C_{1}) + \tau(C_{1}), \text{Int}(C_{2}) + \tau(C_{2}))$$

where $\tau(C) = k/|C|$



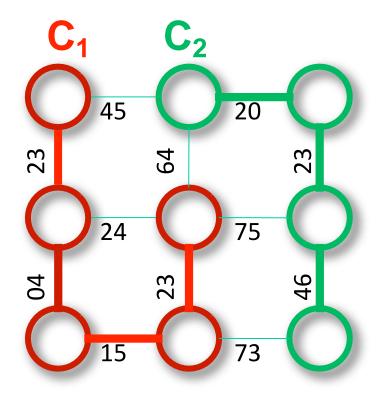
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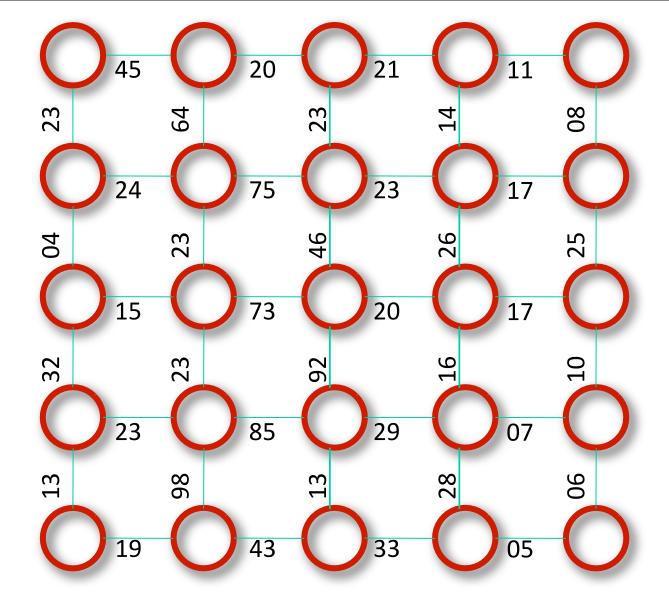
Dif
$$(C_{1,}C_{2}) < \min(\text{Int}(C_{1}) + \tau(C_{1}), \text{Int}(C_{2}) + \tau(C_{2}))$$

where $\tau(C) = k/|C|$

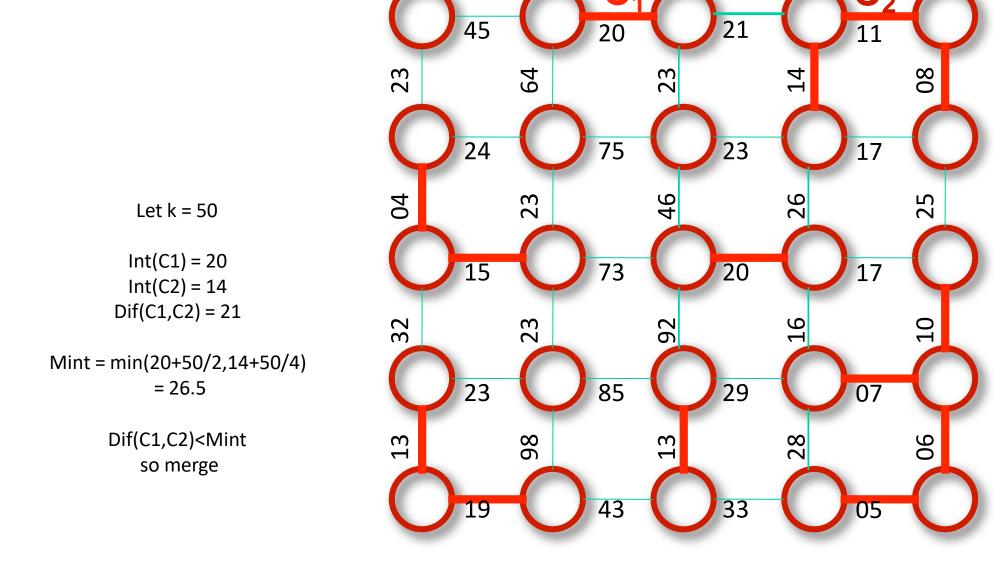


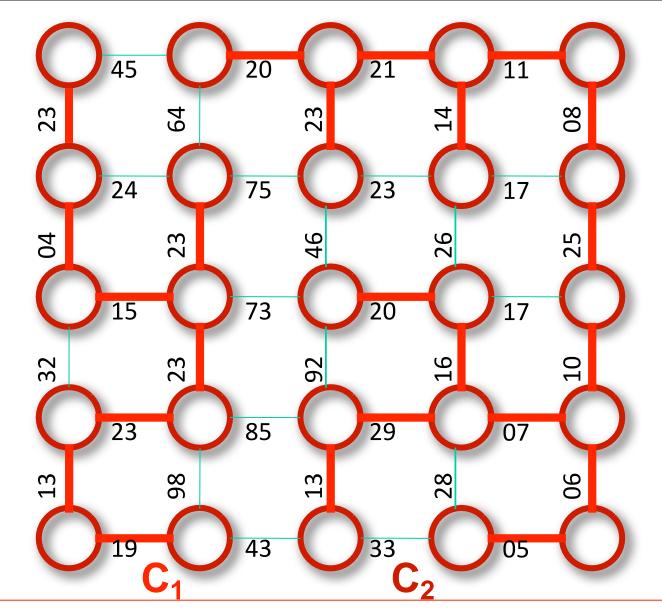
Here:
$$Int(C1) = 23$$

 $Int(C2) = 46$
 $Dif(C1,C2) = min(45,64,73,75)=45$



Sort edge strengths (remembering where they came from)





Let k = 50

Int(C1) = 23Int(C2) = 29

Dif(C1,C2) = 43

Mint = min(23+50/9,19+50/16) = 28.566

Dif(C1,C2)>Mint so don't merge

Example Results

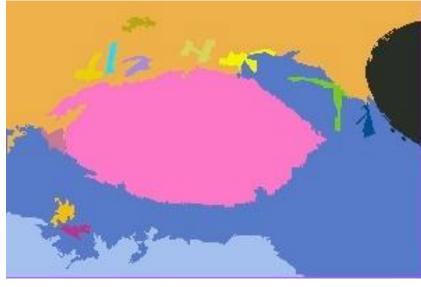


Example Results

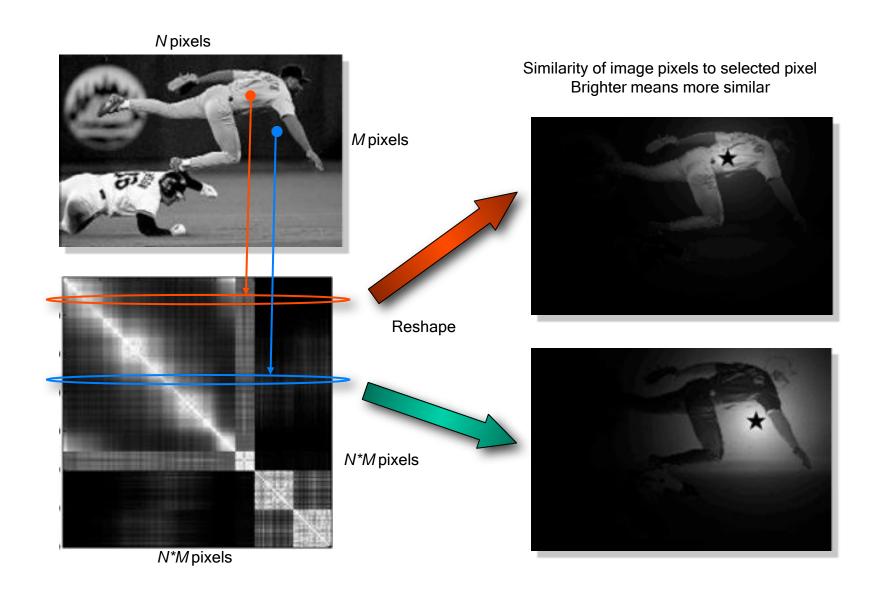


- Very fast
- But sensitive to noise
- Greedily chooses (too) large regions





Affinity Matrix

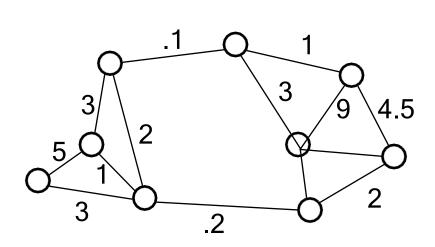


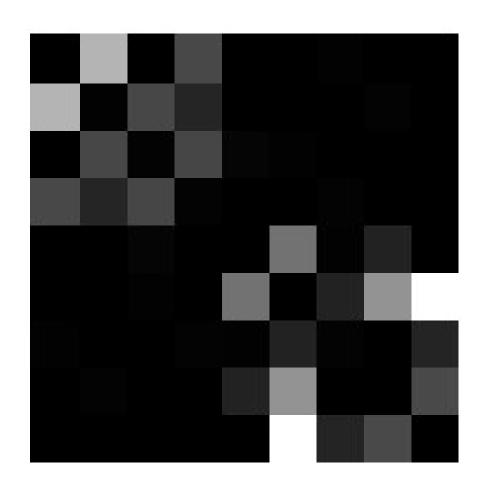
Pixel Similarity Functions

Intensity
$$W(i,j) = e^{\frac{-\left\|I_{(i)} - I_{(j)}\right\|_{2}^{2}}{\sigma_{I}^{2}}}$$

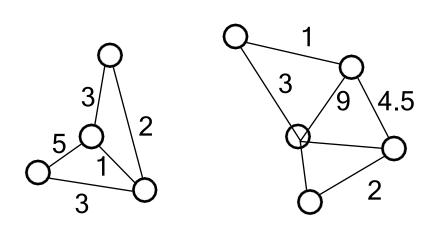
$$W(i,j) = e^{\frac{-\left\|X_{(i)} - X_{(j)}\right\|_{2}^{2}}{\sigma_{X}^{2}}}$$
Texture
$$W(i,j) = e^{\frac{-\left\|c_{(i)} - c_{(j)}\right\|_{2}^{2}}{\sigma_{c}^{2}}}$$

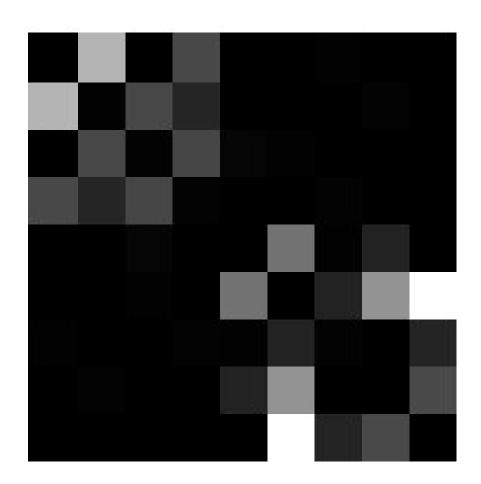
Minimum Cut and Clustering





Minimum Cut and Clustering





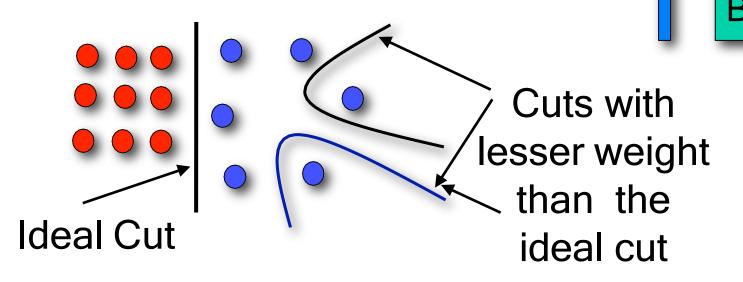
Minimum cut

number of edges in the cut.

Criterion for partition:

$$\min cut(A, B) = \min_{A, B} w(u, v)$$

Problem
Weight of cut is directly proportional to the



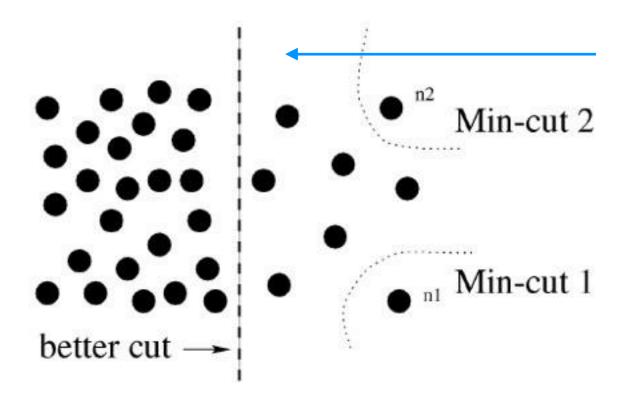
First proposed by Wu and Leahy

Normalized Cut

Normalized cut or balanced cut:

$$N cut(A, B) := cut(A, B)$$

$$\frac{1}{vol(A)} + \frac{1}{vol(B)}$$



finds better cut

Analyzing Normalized Cuts

- State of the art results
- Huge storage requirement and time complexity
- Bias towards partitioning into equal segments
- Has problems with textured backgrounds