



Master in Computer Vision *Barcelona*

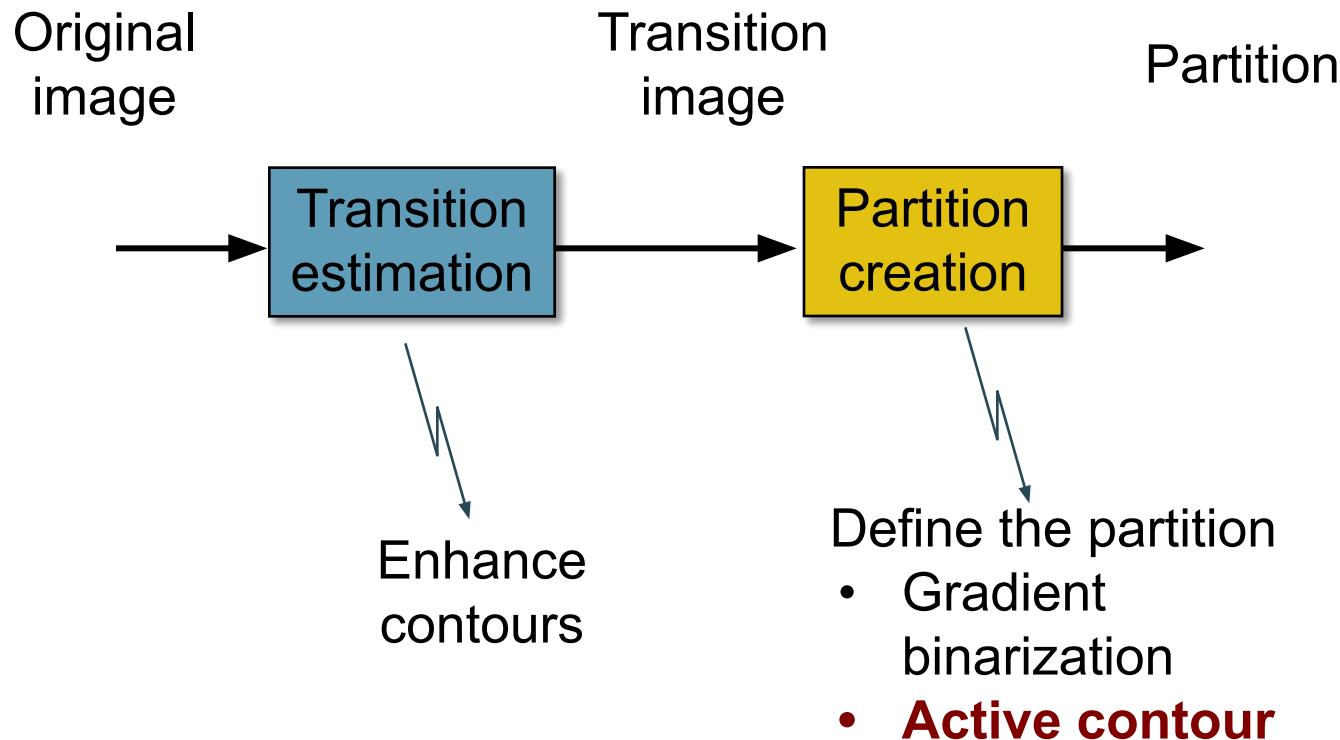
Module: Introduction to human and computer vision

Lecture 10: Grouping, segmentation and classification (II)

Lecturer: Ramon Morros

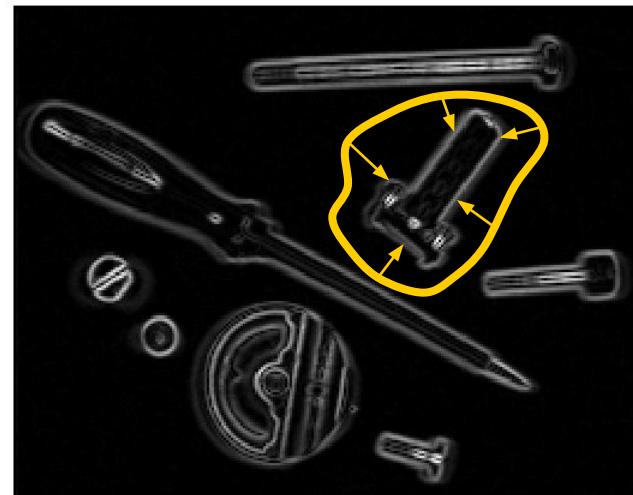
SEGMENTATION

Segmentation: Transition based



Transition based – Active contours

- Robust strategy: Active contours (Snakes)
 - **Evolution of a closed curve towards the points of high**

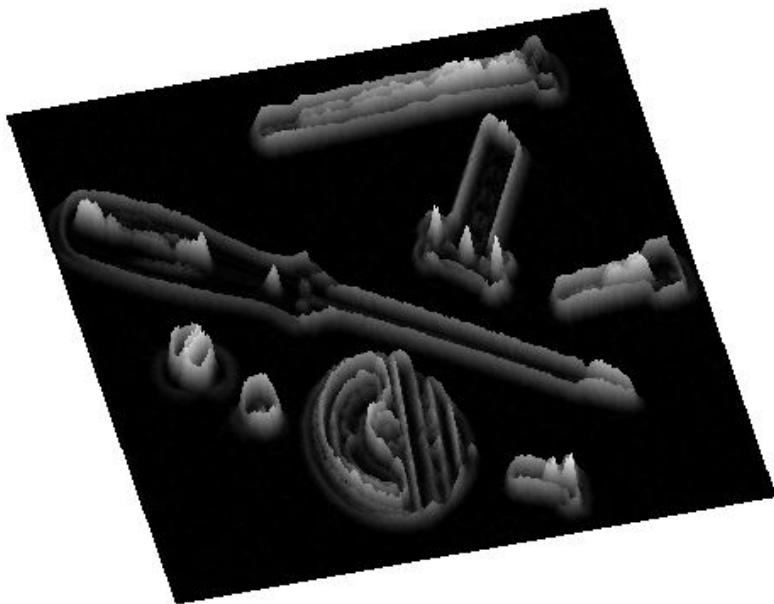
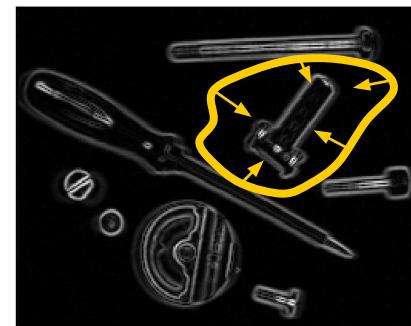


- How to define the curve evolution?
- How to implement it?
- How to define the initial curve?

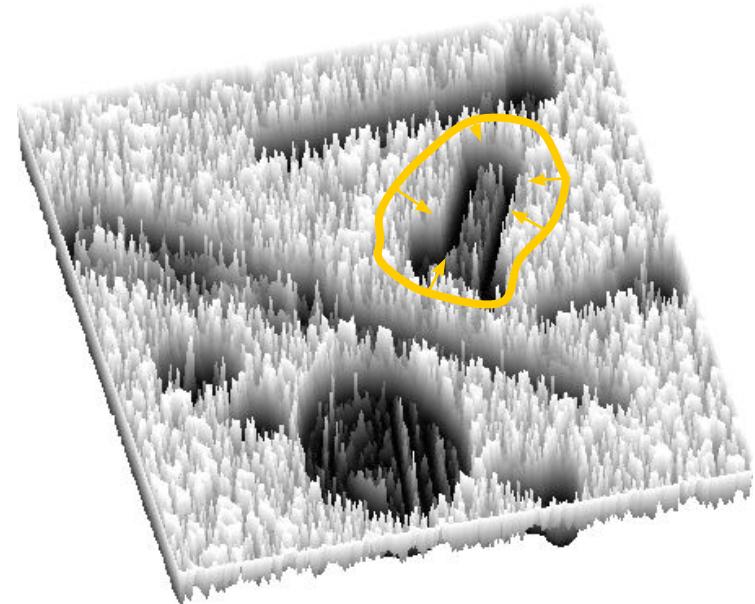
Transition based – Active contours

Gradient:

Find the curve $C(s)$ such that
 $\oint g(\nabla x) ds$ is minimum



3D view of ∇x



3D view of $g(\nabla x)$

Transition based – Active contours

- How to find the minimum of a function?:
 - Compute the gradient of the function:

$$\text{Minimum of } E(i) \Rightarrow \frac{dE(i)}{di} = 0$$

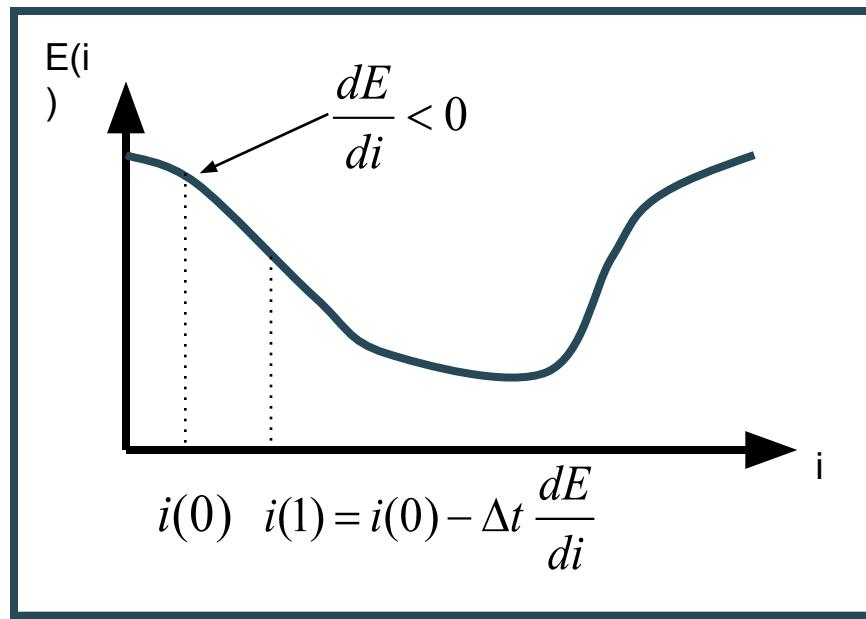
- Gradient descent strategy: iterative search of $i \rightarrow i(t)$. It leads to the following PDE:

$$\frac{\partial i(t)}{\partial t} = -\frac{\partial E(i)}{\partial i}$$

In practice:

$$i(t) - i(t-1) \propto -\frac{\partial E(i)}{\partial i}$$

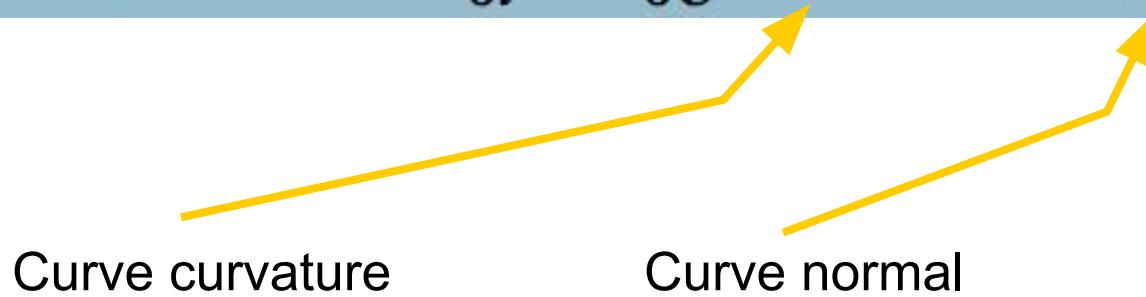
$$i(t) = i(t-1) - \mu \frac{\partial E(i)}{\partial i}$$



Transition based – Active contours

- Gradient descent strategy: iterative search of $i \rightarrow i(t)$.
It leads to:

$$E(C) = \oint g(\nabla x) ds \Rightarrow \frac{\partial C}{\partial t} = -\frac{\partial E}{\partial C} = gk\vec{N} - \langle \nabla g, \vec{N} \rangle \vec{N}$$



Transition based – Active contours

Brief reminder on differential geometry

- Continuous image model: $i, j \in \mathbb{R}$
- Curve parametrization:

$$C(p) = \{(i(p), j(p)), p \in [a, b]\}$$

- Arc-length parametrization:

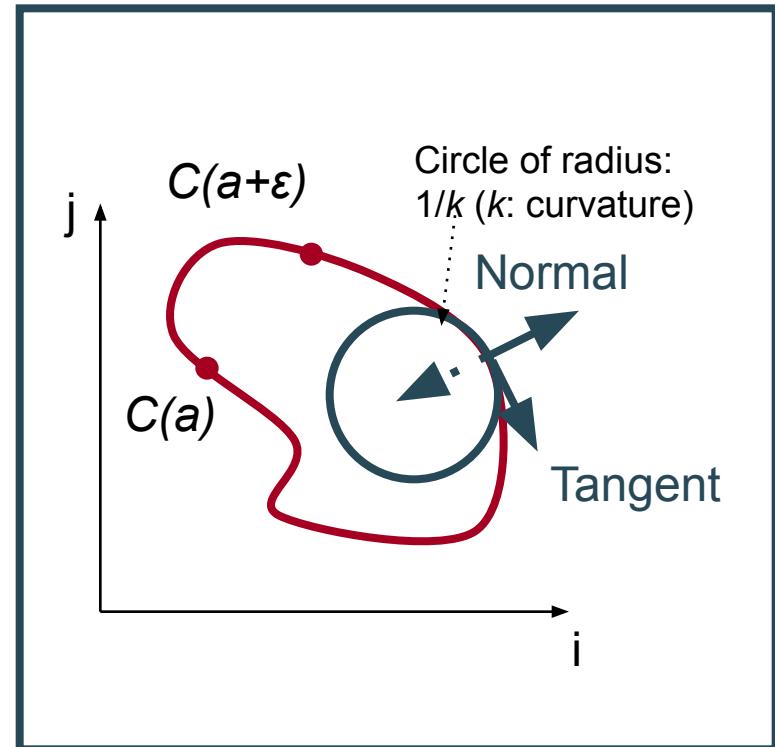
$$s \text{ such that } \left\| \frac{dC}{ds} \right\| = 1$$

$$C(s) = \{(i(s), j(s)), s \in [0, \text{Length}]\}$$

- Tangent and Normal:

$$\vec{T} = \frac{dC}{ds} = \left(\frac{di(s)}{ds}, \frac{dj(s)}{ds} \right)$$

$$k\vec{N} = \frac{d^2C}{ds^2}$$



$$\begin{aligned} \left\langle \frac{dC}{ds}, \frac{dC}{ds} \right\rangle = 1 &\Rightarrow \left\langle \frac{dC}{ds}, \frac{d^2C}{ds^2} \right\rangle = 0 \\ \vec{T} \perp \frac{d^2C}{ds^2} &= k\vec{N} \end{aligned}$$

Transition based – Active contours

- Using as criterion the geodesic length:

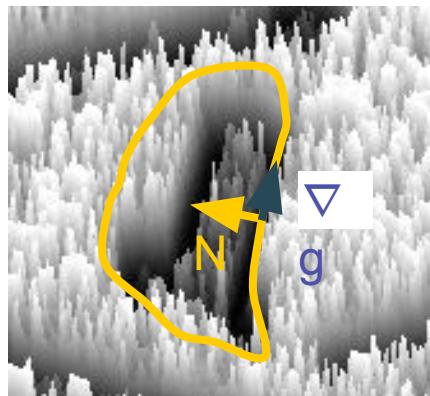
$$E(C) = \oint g(\nabla x) ds \Rightarrow \frac{\partial C}{\partial t} = -\frac{\partial E}{\partial C} = gk\vec{N} - \langle \nabla g, \vec{N} \rangle \vec{N}$$

Internal force:

- Curvature motion
- Smoothing

External force:

- Force the curve evolution towards minima of g
- The evolution stops when the gradient of g is perpendicular to the curve normal



Note: Other (but similar) expressions are possible for the internal and external forces

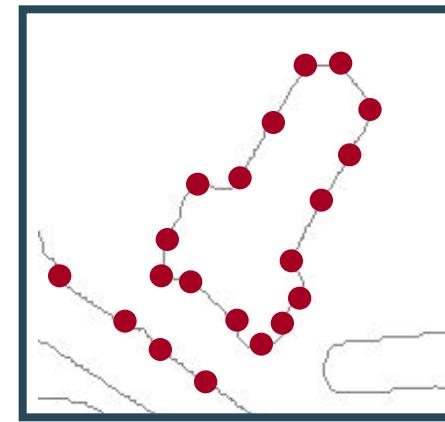
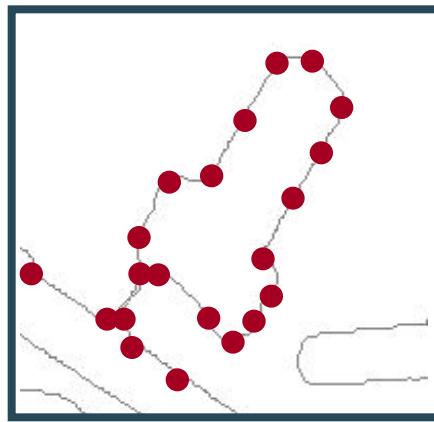
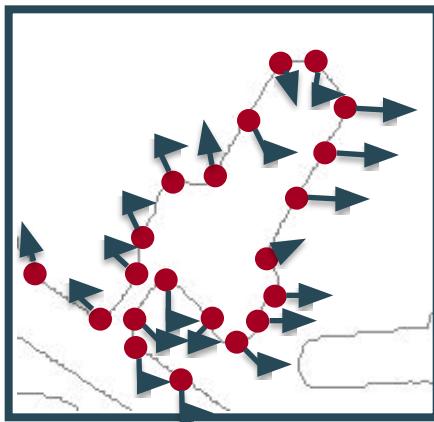
In practice:

$$C(t) = C(t-1) + \mu(gk - \langle \nabla g, \vec{N} \rangle) \vec{N}$$

Transition based – Active contours

- Curve defined by control points:
 - Displace the control point following:
 - Resample the curve when it shrinks
 - Handle topological changes
- Other implementations are possible (for example: level set)

$$(gk - \langle \nabla g, \vec{N} \rangle) \vec{N}$$



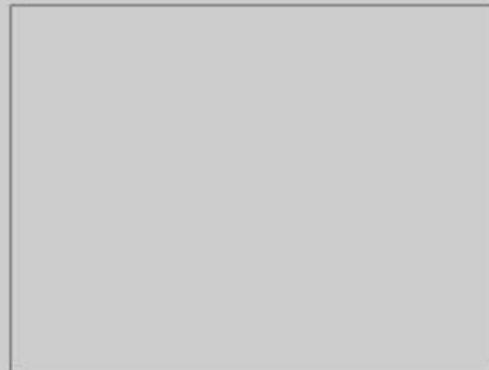
Transition based – Active contours



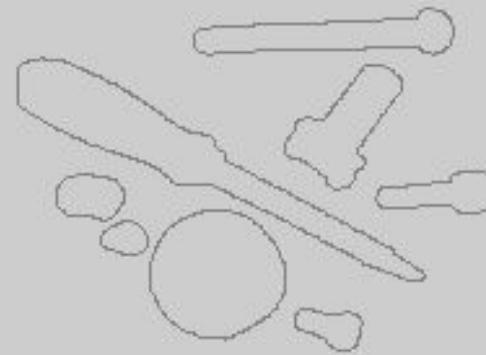
Original image x



$G(\nabla x)$



Initial contour

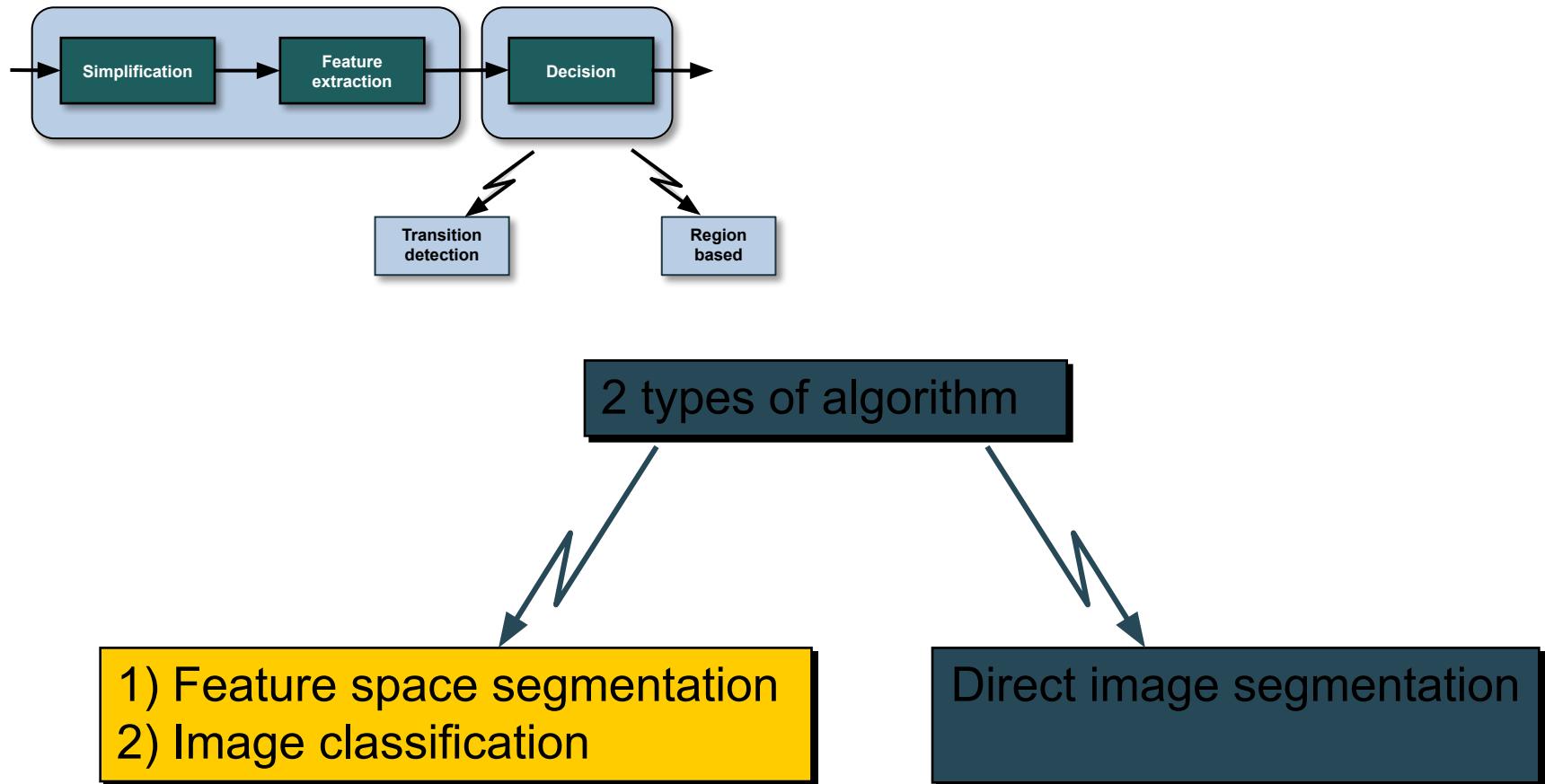


Final contour

Segmentation: Transition based

- 3 main steps for segmentation:
 - Simplification, feature extraction and decision
 - Decision can be based on discontinuity (transition-based) or homogeneity (region-based)
- Transition-based segmentation:
 - Estimate transitions: gradient (may be linear or not)
 - Decision:
 - **Thresholding:** difficult because of threshold definition, noise and open contour
 - **Active contours:**
 - Curve evolution to minimize the length “weighted” by the gradient
 - Gradient descent leads to a combination of internal (smoothing) forces and external force (attraction to data)

Segmentation: Homogeneity based



Homogeneity based – Feature space segmentation

Approach:

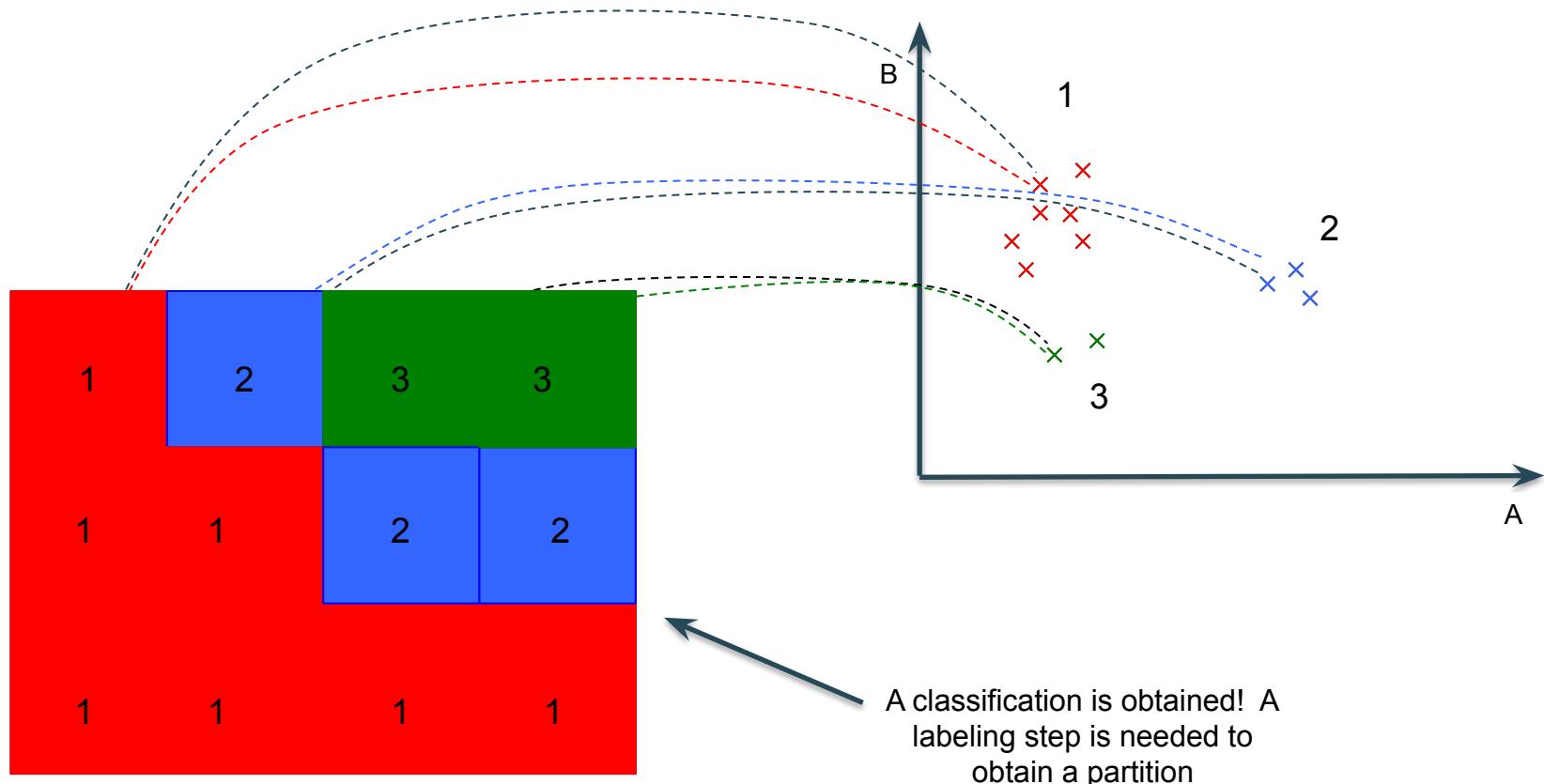
- Study the feature space
- Defines classes in the feature space
(feature space segmentation)
- Region definition
(image pixel classification)

Examples:

- Mono-dimensional feature space (gray level)
 - 2 classes
 - N classes
- Multidimensional feature space

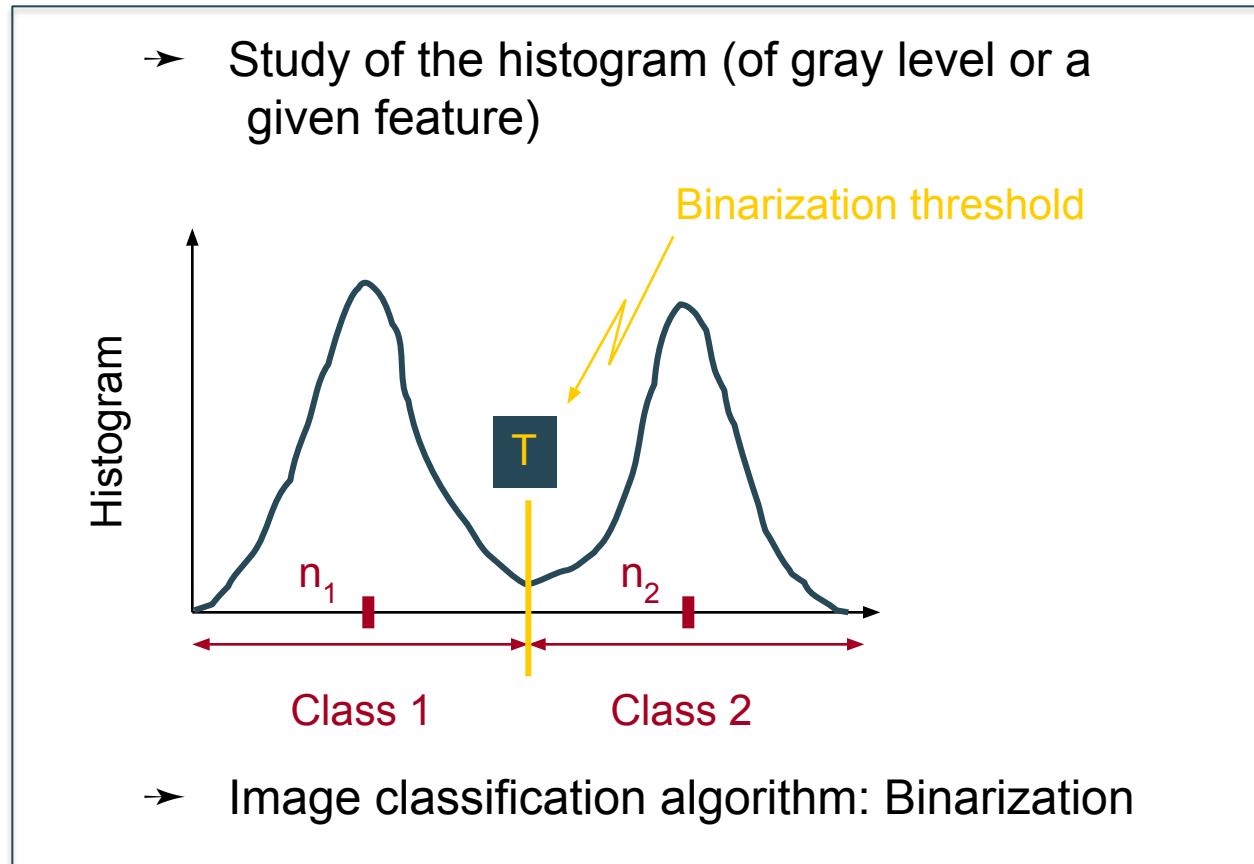
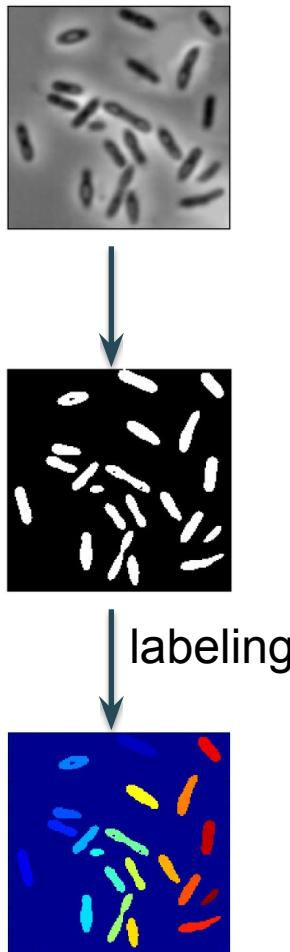
Homogeneity based – Feature space segmentation

Example: bi-dimensional feature space



Homogeneity based – Feature space segmentation

Mono-dimensional feature space, 2 classes



Homogeneity based – Feature space segmentation

Mono-dimensional feature space, 2 classes: threshold

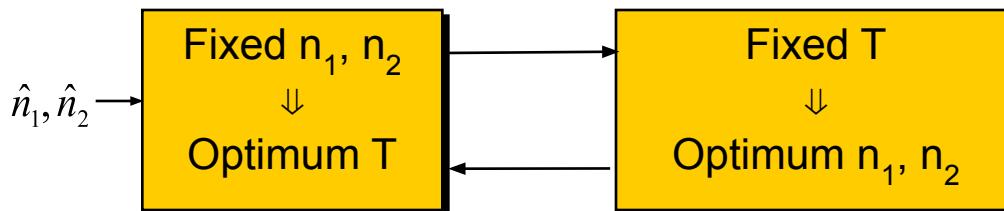
- If the class centers n_1 y n_2 are known (e.g.: histogram maxima)

- Criterion: Intra class minimum variance

$$C = \operatorname{Min}_T \left[\sum_{0 \leq n \leq T} (n - n_1)^2 h(n) + \sum_{T < n \leq M} (n - n_2)^2 h(n) \right]$$

- If class centers are NOT known

→ Iterative estimation



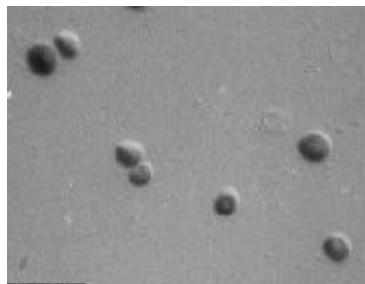
→ Local minimum

$$\begin{aligned} n_1 &= \frac{\sum_{0 \leq n \leq T} nh(n)}{\sum_{0 \leq n \leq T} h(n)} \\ n_2 &= \frac{\sum_{T < n \leq M} nh(n)}{\sum_{T < n \leq M} h(n)} \end{aligned}$$

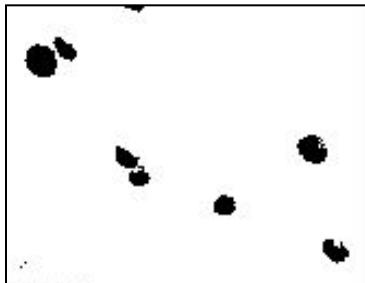
Mean of
each class

Homogeneity based – Feature space segmentation

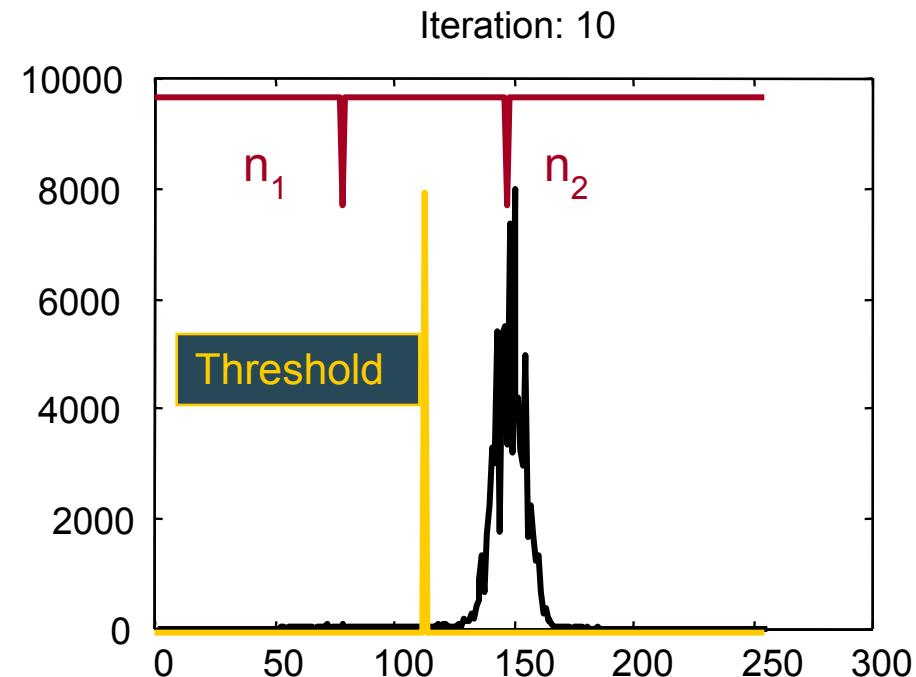
Mono-dimensional feature space, 2 classes: threshold



Original



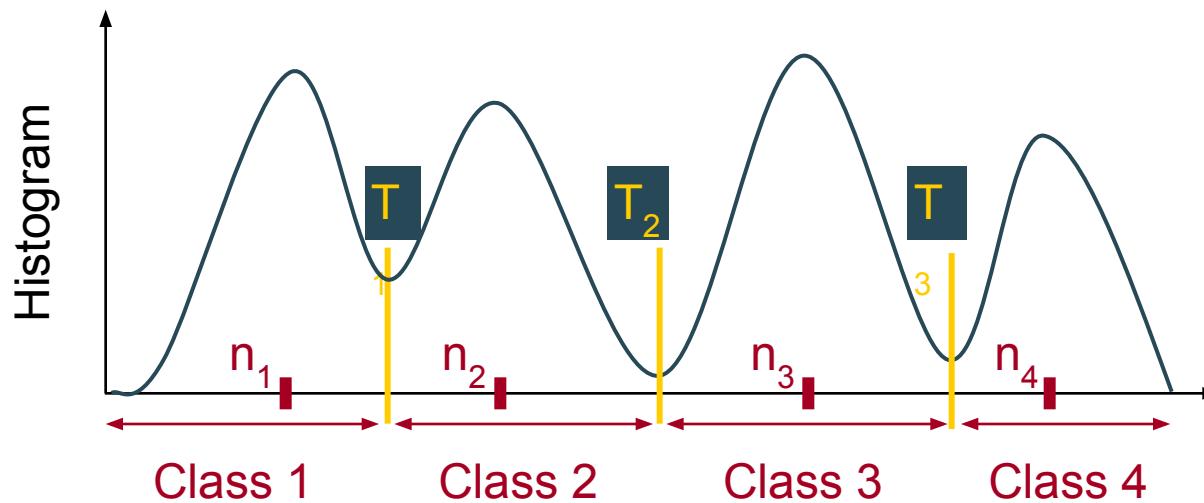
Binarized image



Homogeneity based – Feature space segmentation – K-Means

Mono-dimensional feature space, K classes

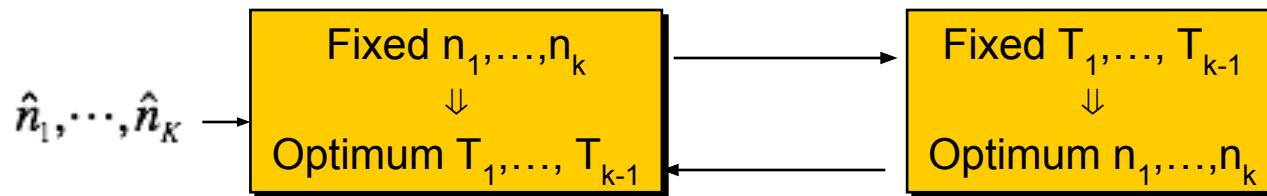
→ Histogram analysis



→ Classification algorithm: “K-means”

Homogeneity based – Feature space segmentation – K-Means

K-Means algorithm



- All T combinations (may be complex!)
- Euclidean distance

Individual class mean

$$T_1 = (n_1 + n_2) / 2$$

...

$$T_{k-1} = (n_{k-2} + n_{k-1}) / 2$$

$$n_1 = \sum_{0 \leq n \leq T_1} nh(n) / \sum_{0 \leq n \leq T_1} h(n)$$

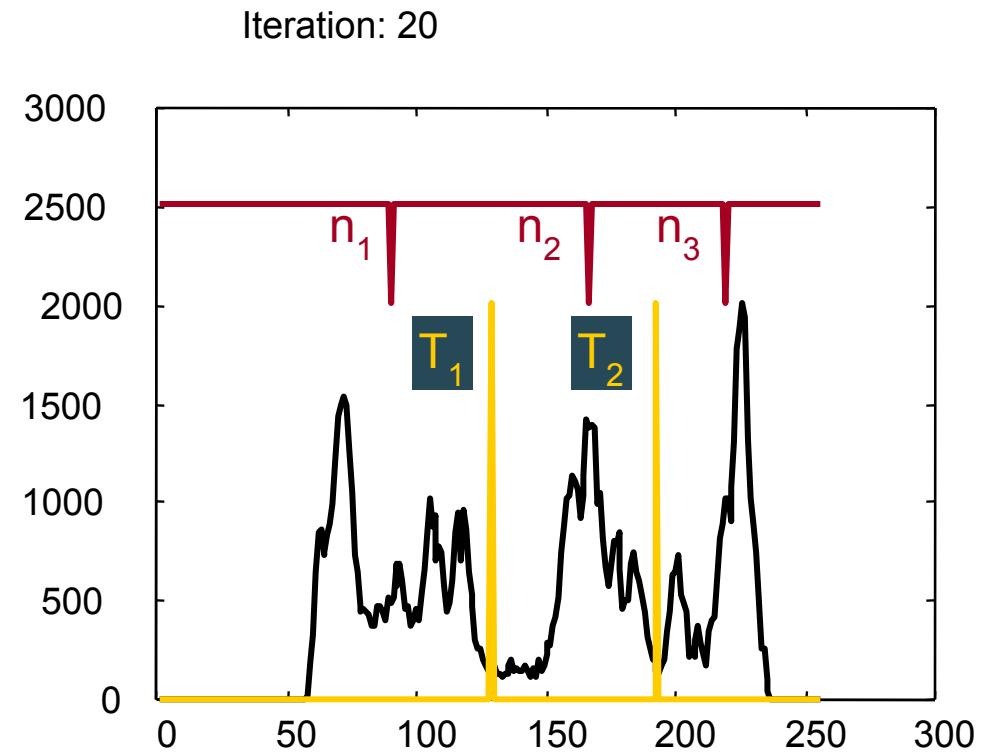
...

$$n_K = \sum_{T_{k-1} < n \leq M} nh(n) / \sum_{T_{k-1} < n \leq M} h(n)$$

→ Local minimum

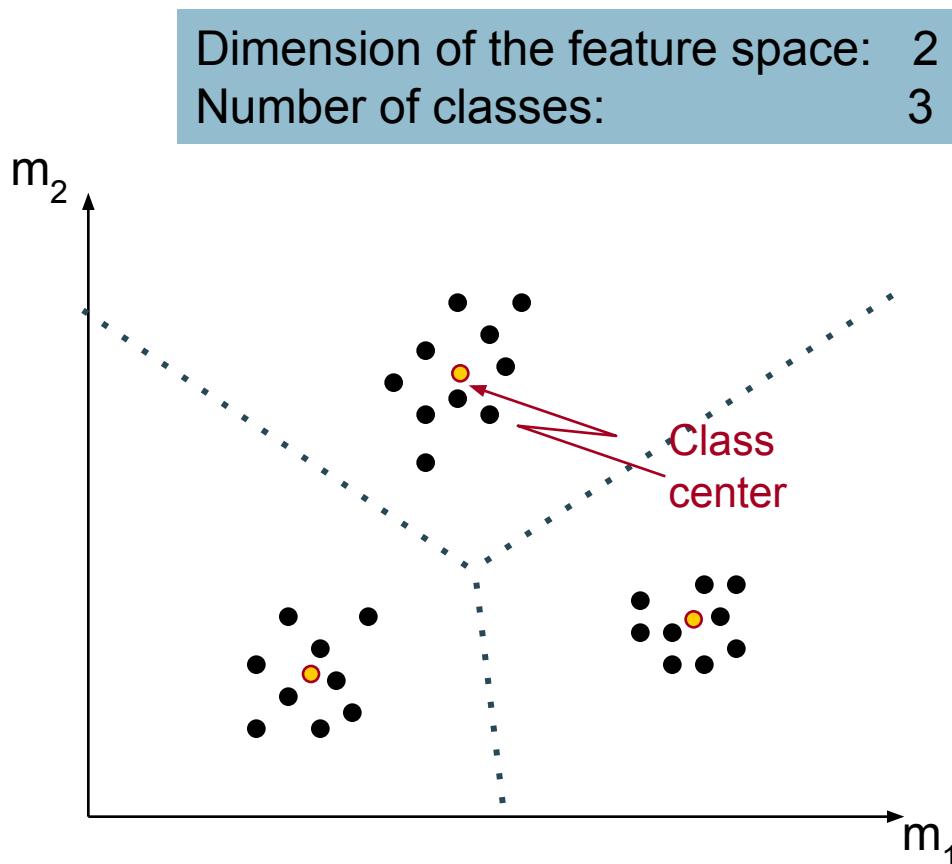
Homogeneity based – Feature space segmentation – K-Means

K-Means example : 3 classes



Homogeneity based – Feature space segmentation – K-Means

Multi-dimensional feature space, K classes



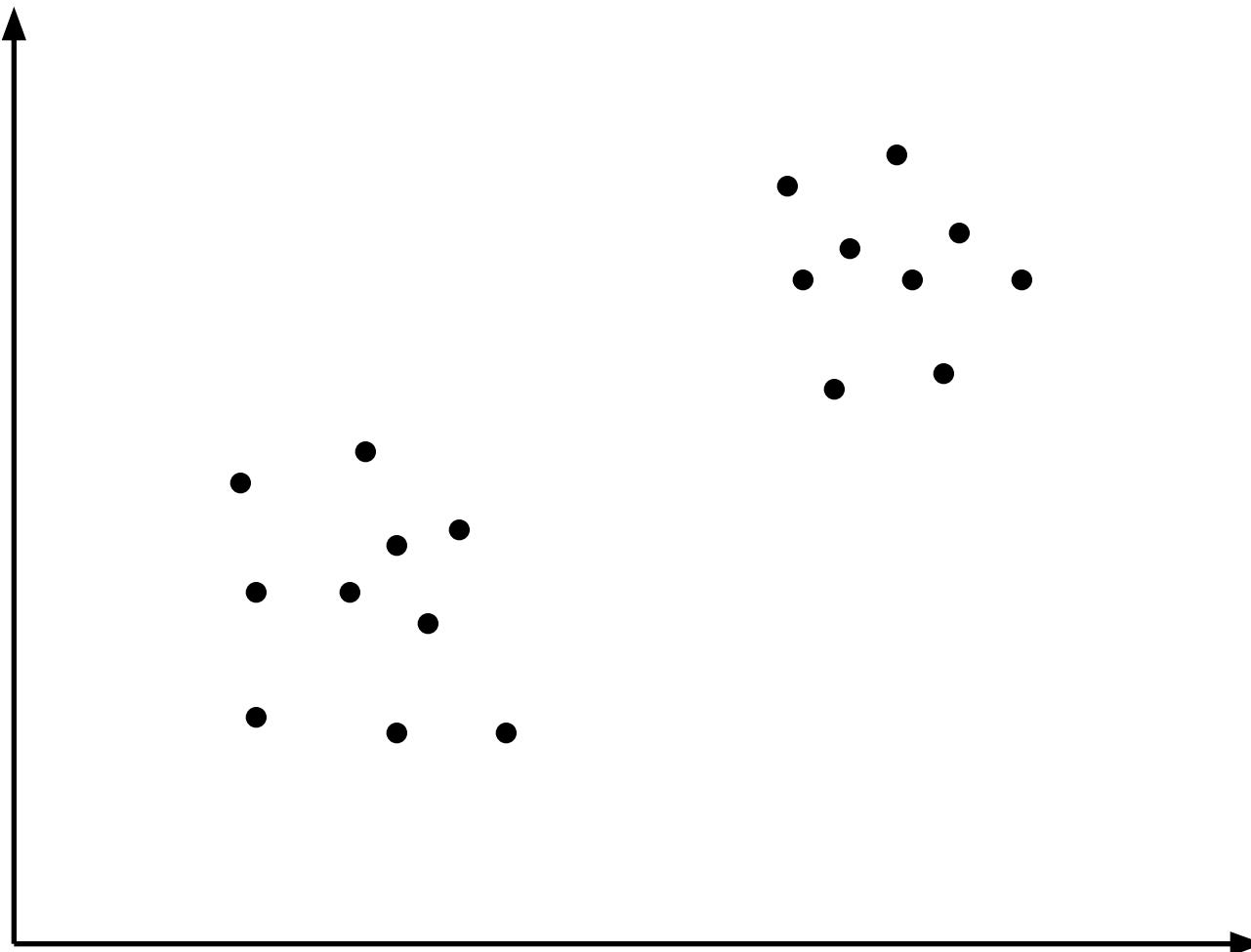
1. Initialize K classes. Compute the centers of each class
2. For each point:
 - a. Compute the distances between the point and the class centers
 - b. Assign the point to the closest class
3. Update the class centers
4. Repeat 2 & 3 until no change (in assignments or center values) is observed.

Max Lloyd algorithm

Complexity: $O(n \cdot k \cdot d \cdot i)$
d: dimension, n: number of points, i: iterations

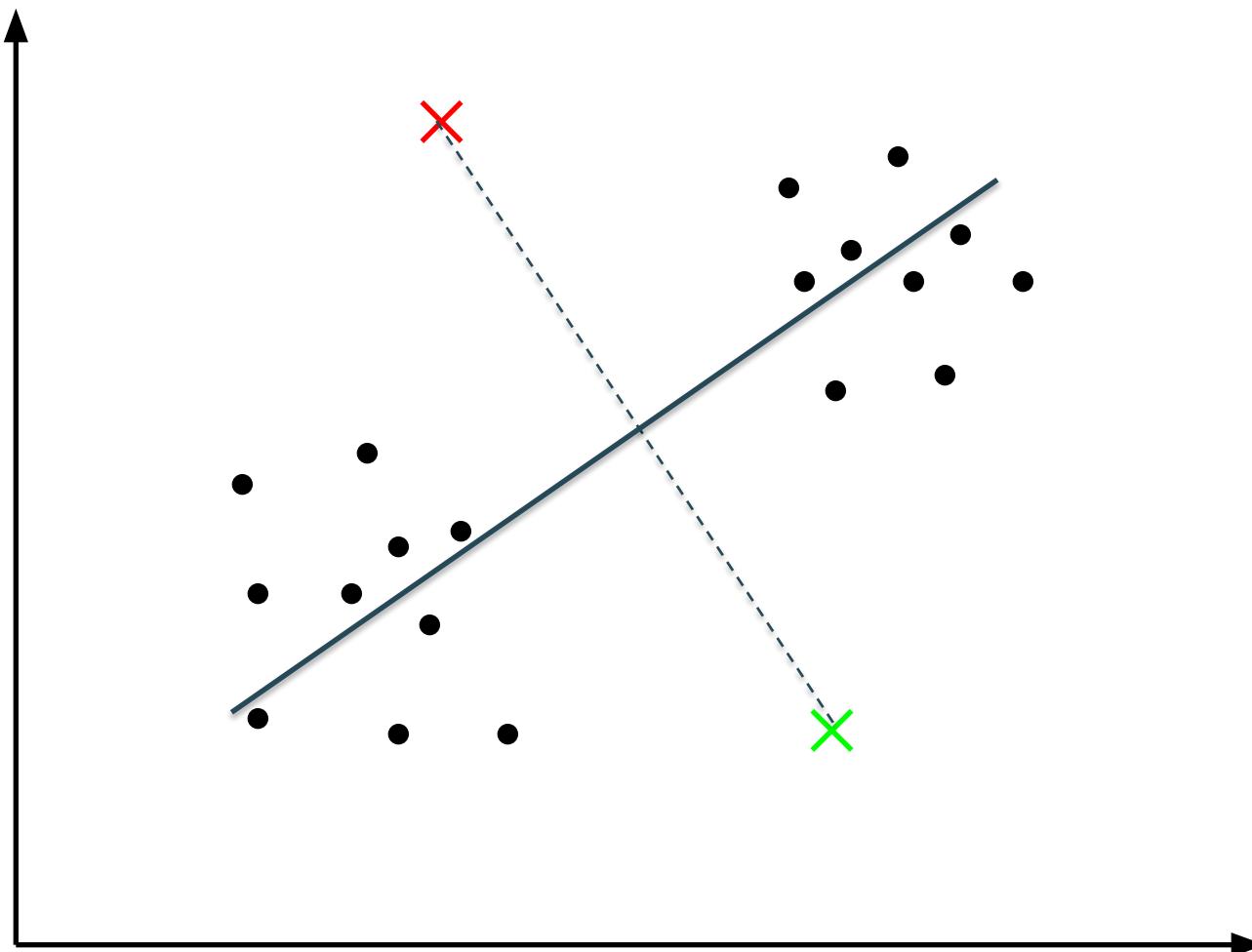
Homogeneity based – Feature space segmentation – K-Means

Example:



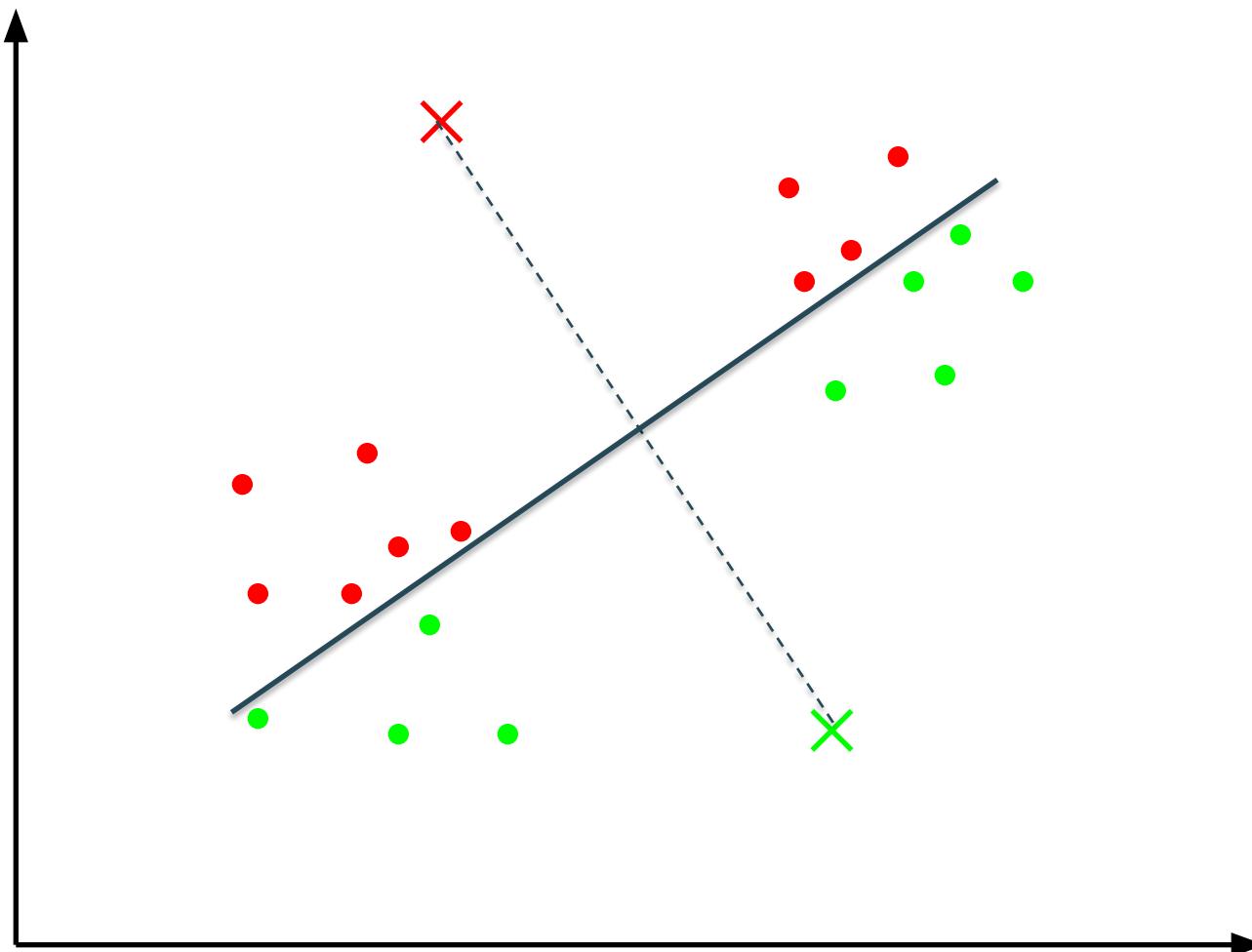
Homogeneity based – Feature space segmentation – K-Means

Example:



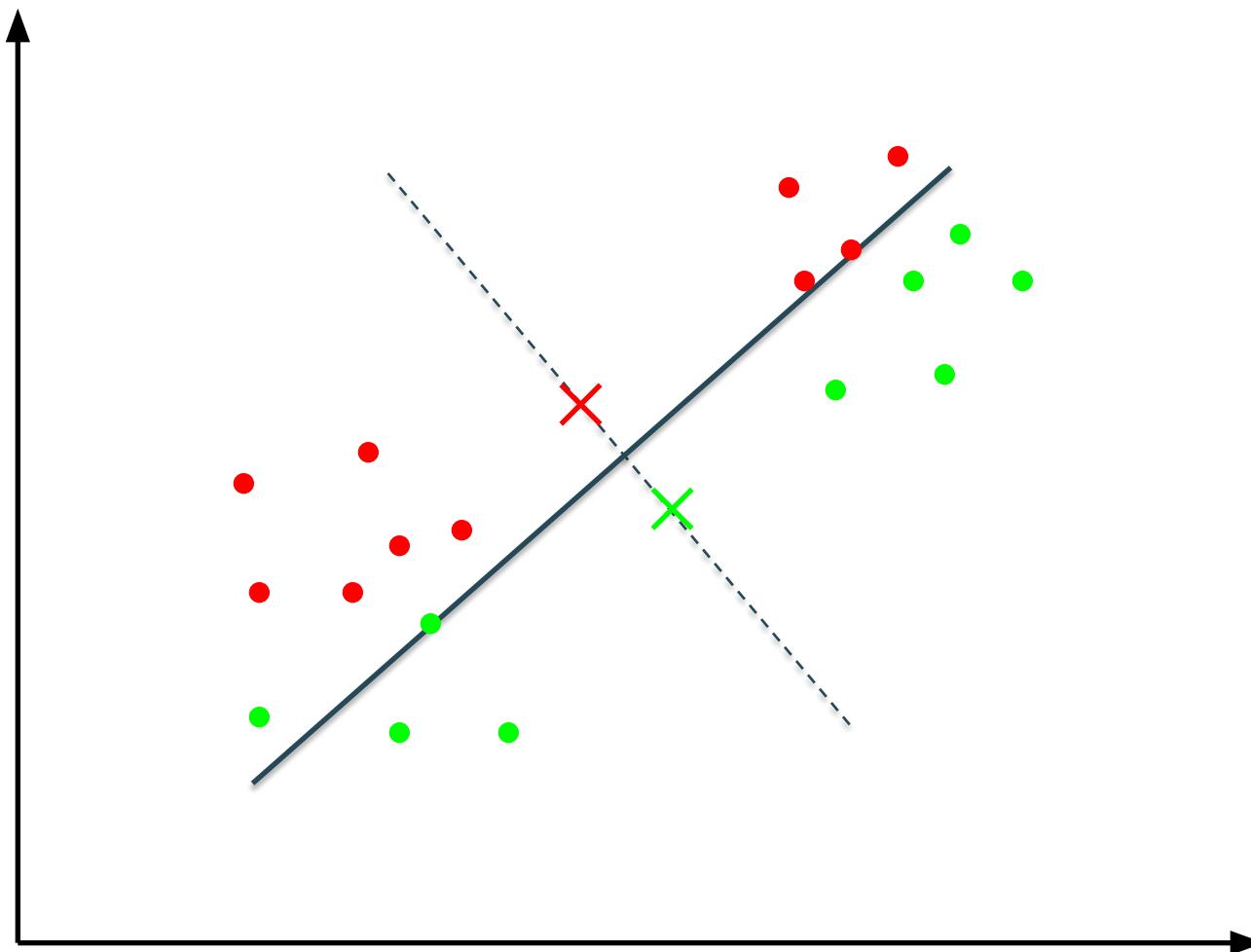
Homogeneity based – Feature space segmentation – K-Means

Example:



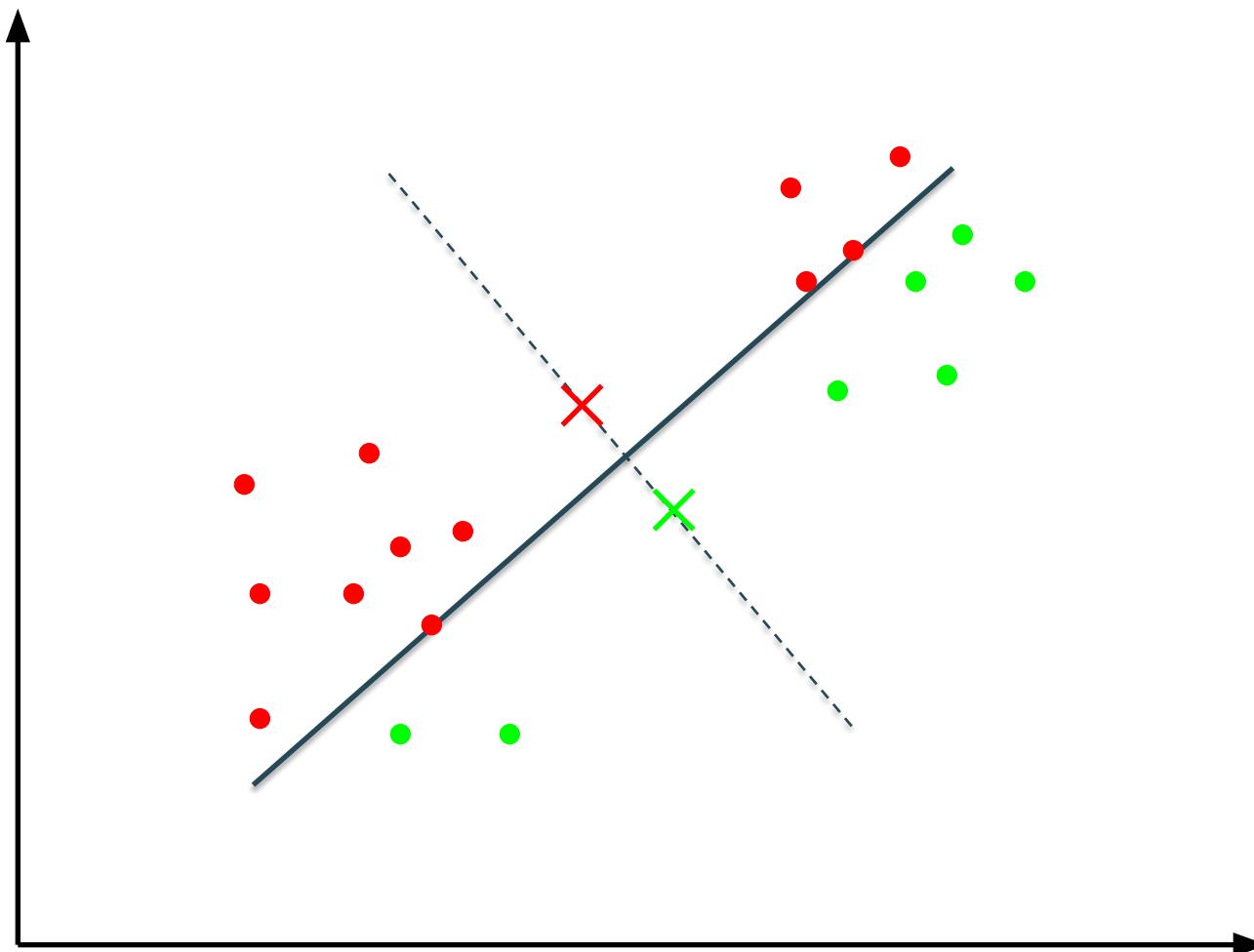
Homogeneity based – Feature space segmentation – K-Means

Example:



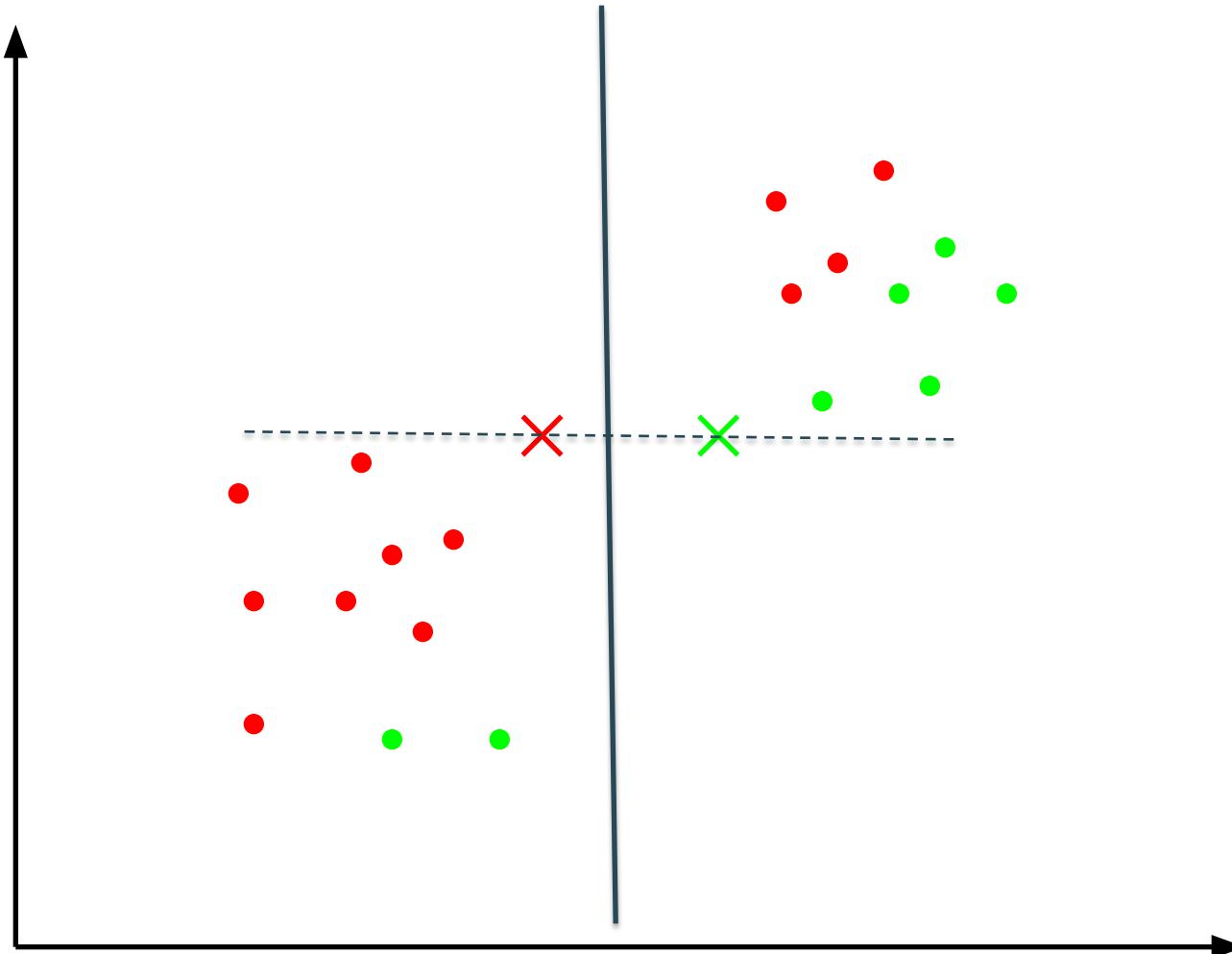
Homogeneity based – Feature space segmentation – K-Means

Example:



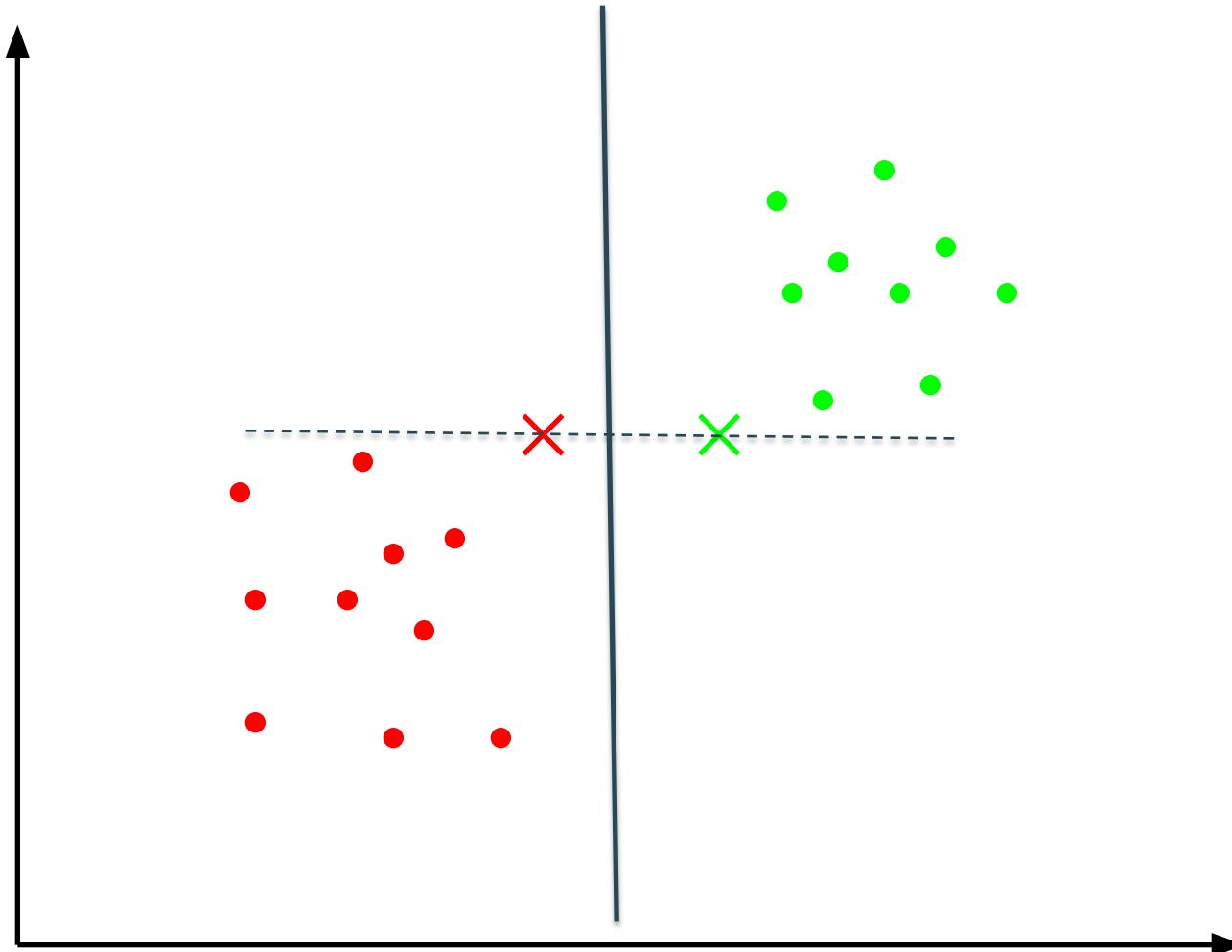
Homogeneity based – Feature space segmentation – K-Means

Example:



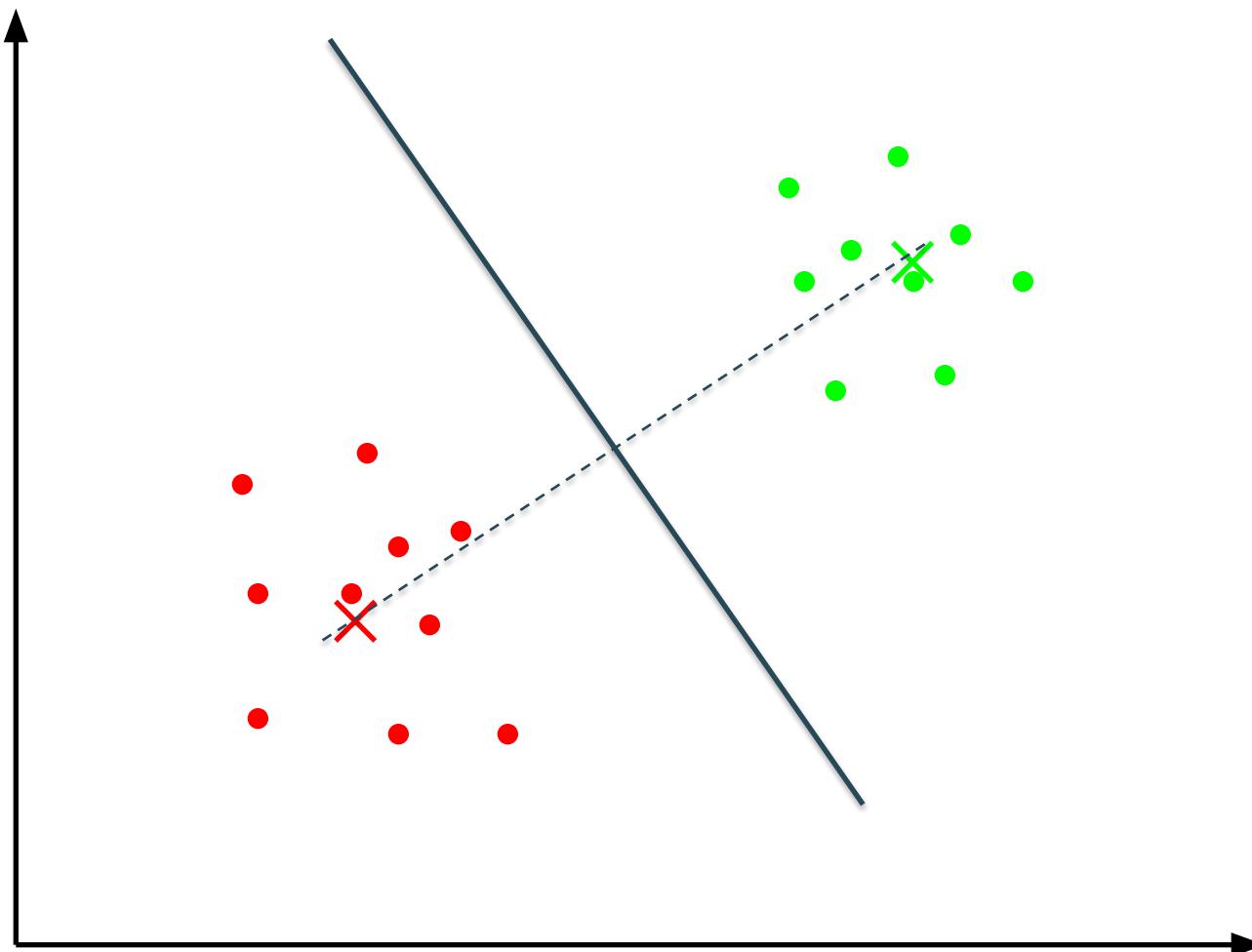
Homogeneity based – Feature space segmentation – K-Means

Example:



Homogeneity based – Feature space segmentation – K-Means

Example:



Homogeneity based – Feature space segmentation – K-Means

- Convergence is guaranteed ... but may be to a local minimum → application optimized strategies can be added
 - The straightforward implementation of such method may be too slow → there are proposed optimizations, especially for the classification step.
 - Other distances are also possible (K-Medoids).
 - Hard assignment of data points to clusters: small shift of a data point can flip it to a different cluster → can be solved by a soft assignment based on a probabilistic approach.

Gaussian Mixture Models

Slide credit: Dr. Antonio M. López (UAB)

Homogeneity based – Feature space segmentation – K-Means

Example: Face (skin) detection

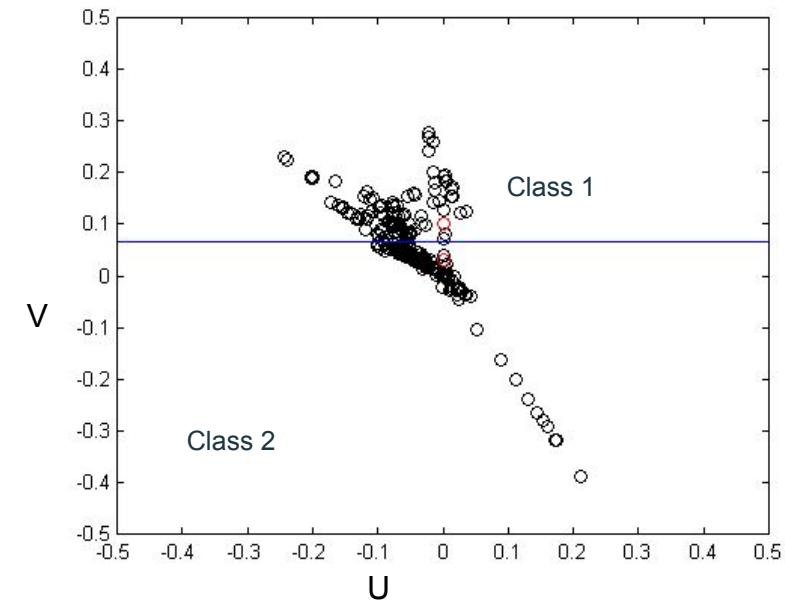


Original
image

$U=B-Y$
component



$V=R-Y$
Component

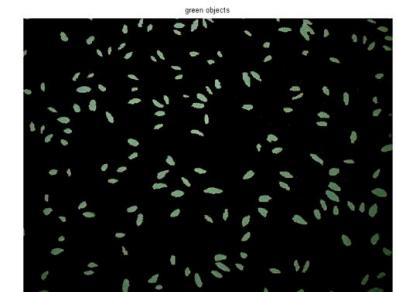
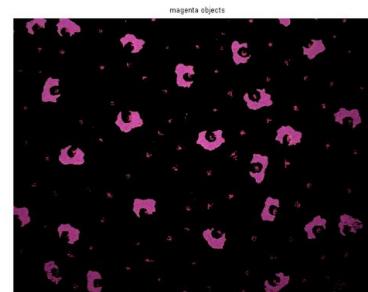
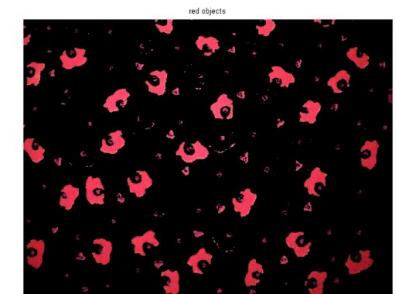
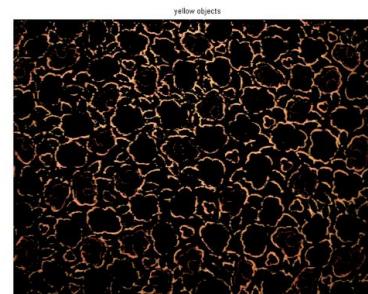
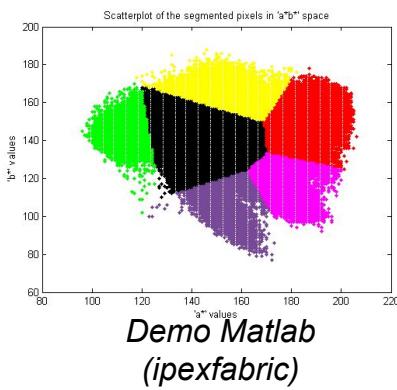
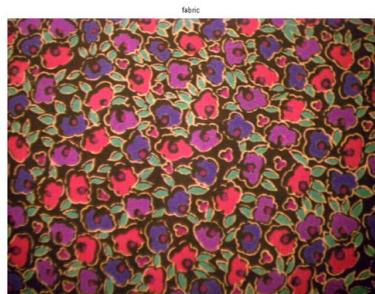


Binarization

Homogeneity based – Feature space segmentation – K-Means

Example: Color segmentation

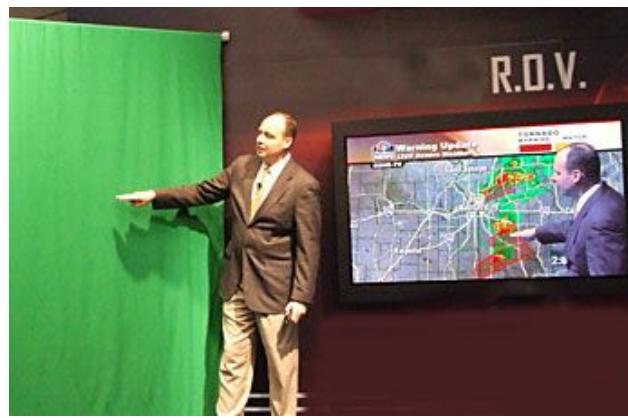
Goal: Segment the regions corresponding to the N dominant color of the image
Color space: CIE Lab



Homogeneity based – Feature space segmentation – K-Means

Example: Chroma Key

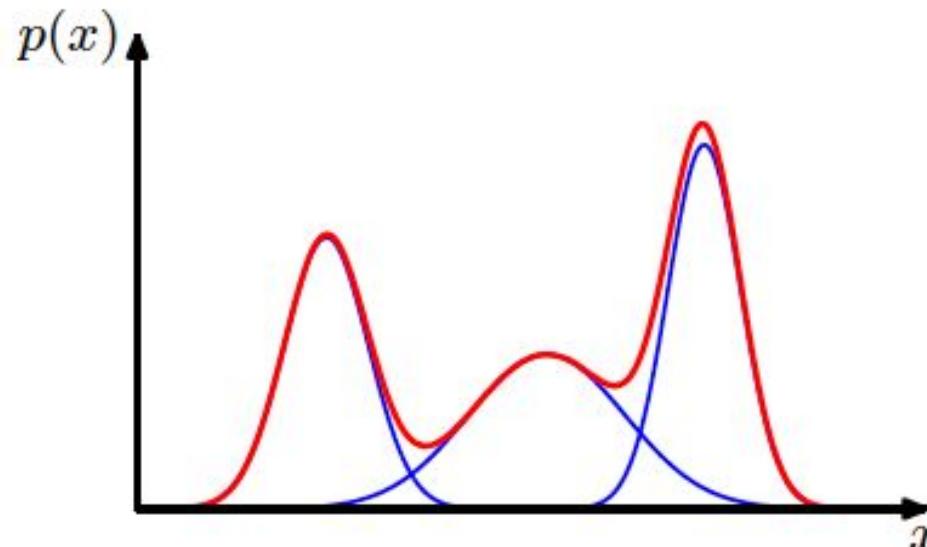
- Used in TV, cine or photography
- Consist in changing the scene background
- Background is generally blue or green. It is segmented and the visible background pixels are substituted by a different background.
- Possible issues: incoherent background/scene illumination or shadows



Homogeneity based – FS segmentation : Mixture Models

- In k-means, pixels are clustered using hard assignments
 - Each pixel goes to closest cluster center
 - A probabilistic approach, where each pixel helps estimating more than one cluster, may be more robust

→ **Probabilistic Mixture Model**



Homogeneity based – FS segmentation : Mixture Models

- The $p(x | k)$ are, in fact, parametric densities: $p(x | \theta_k)$
- A widely used approach is to assume that all the $p(x | k)$ from the same family of densities
- In the limit, even smooth densities can be approximated
- Multivariate gaussians are a very popular choice
- Applications:
 - Probability density estimation
 - **Clustering: every component is a cluster center**

Homogeneity based – FS segmentation : GMM

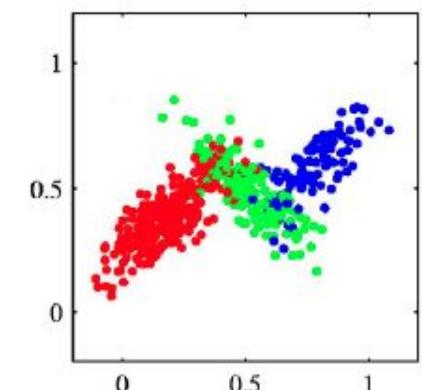
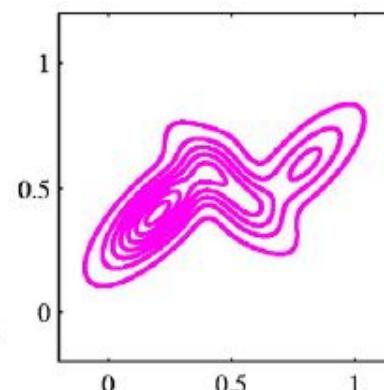
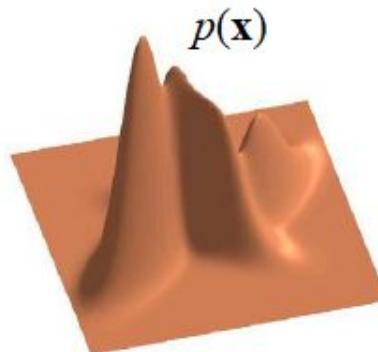
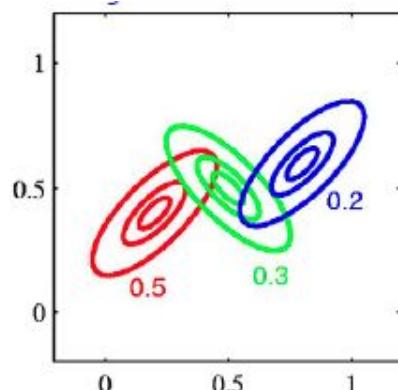
Good choice for the components → Gaussians

$$p(x | \theta) = \sum_{k=1}^K \pi_k p(x | \theta_k)$$

$$0 \leq \pi_k \leq 1, \quad \sum_{k=1}^K \pi_k = 1$$

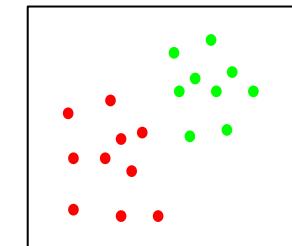
$$p(x | \theta_k) = N(x | \mu_k, \Sigma_k) \propto \exp\left(\frac{-(x - \mu_k)^T \Sigma_k^{-1} (x - \mu_k)}{2}\right)$$

Gaussians can be seen as oriented “blobs” in feature space



Homogeneity based – FS segmentation : GMM

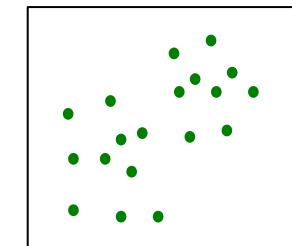
- Mixture parameter estimation:
 - Labeled data: we know which component generates each Gaussian → Easy!!



$$\pi_k = \frac{n_k}{N}, \quad \mu_k = \frac{1}{n_k} \sum_{i|l_i=k} x_i, \quad \Sigma_k = \frac{1}{n_k} \sum_{i|l_i=k} (x_i - \mu_k)(x_i - \mu_k)^T$$

Maximum Likelihood (ML) estimates

- Unlabeled data: the assignments are not known
 - Guessed based on the current mixture distribution estimate
 - Soft assignments (posterior probabilities)



Homogeneity based – FS segmentation : GMM

- To maximize $\log(L(\theta | x))$, we can derive with respect to the parameters and equal to zero. We obtain:

$$\mu_k = \frac{1}{N_k} \sum_{i=1}^N \gamma_k(x_i) \cdot x_i$$

$$N_k = \sum_{i=1}^N \gamma_k(x_i)$$

$$\Sigma_k = \frac{1}{N_k} \sum_{i=1}^N \gamma_k(x_i) \cdot (x_i - \mu_k) \cdot (x_i - \mu_k)^T$$

$$\pi_k = \frac{N_k}{N}$$

$$\gamma_k(x_i) = \frac{\pi_k N(x_i | \mu_k, \Sigma_k)}{\sum_{j=1}^K \pi_j N(x_i | \mu_j, \Sigma_j)}$$

Posterior probability of k given x_i , or in other words, responsibility that the Gaussian k takes for explaining the observation x_i .

Slide credit: Dr. Antonio M. López



Homogeneity based – FS segmentation : GMM

- $\gamma_k(x_i)$ represents the posterior prob. of k given x_i
→ responsibility that gaussian k takes for explaining observation x_i

$$p(x | k) \sim N(x | \mu_k, \Sigma_k)$$
$$p(k | x) = \frac{P(k)p(x | k)}{p(x)} = \frac{\pi_k N(x | \mu_k, \Sigma_k)}{\sum_{j=1}^K \pi_j N(x | \mu_j, \Sigma_j)} \rightarrow \gamma_k(x_i) = p(k | x_i)$$
$$p(x) = \sum_{k=1}^K \pi_k p(x | \theta_k)$$

Slide credit: Dr. Antonio M. López

Homogeneity based – FS segmentation : GMM

Interpretation

$$\gamma_k(x_i) = \frac{\pi_k N(x_i | \mu_k, \Sigma_k)}{\sum_{j=1}^K \pi_j N(x_i | \mu_j, \Sigma_j)}$$

Notice: if **K=1** then $\gamma_1(x_n)=1$.

This corresponds to the ML estimation of the parameters of a single Gaussian kernel.

$$\left. \begin{array}{l} \mu_k = \frac{1}{N_k} \sum_{i=1}^N \gamma_k(x_i) \cdot x_i \\ N_k = \sum_{i=1}^N \gamma_k(x_i) \\ \Sigma_k = \frac{1}{N_k} \sum_{i=1}^N \gamma_k(x_i) \cdot (x_i - \mu_k) \cdot (x_i - \mu_k)^T \\ \pi_k = \frac{N_k}{N} \end{array} \right\}$$

Weighted mean of all the points in the data set

Effective number of points assigned to cluster **k**

Covariance as weighted variances of the data wrt to each component estimated mean

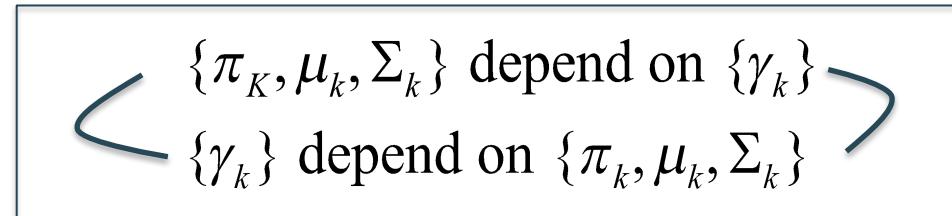
Average responsibility that each component takes for explaining the data

Slide credit: Dr. Antonio M. López



Homogeneity based – FS segmentation : GMM

- If $K > 1$, there is no closed form solution

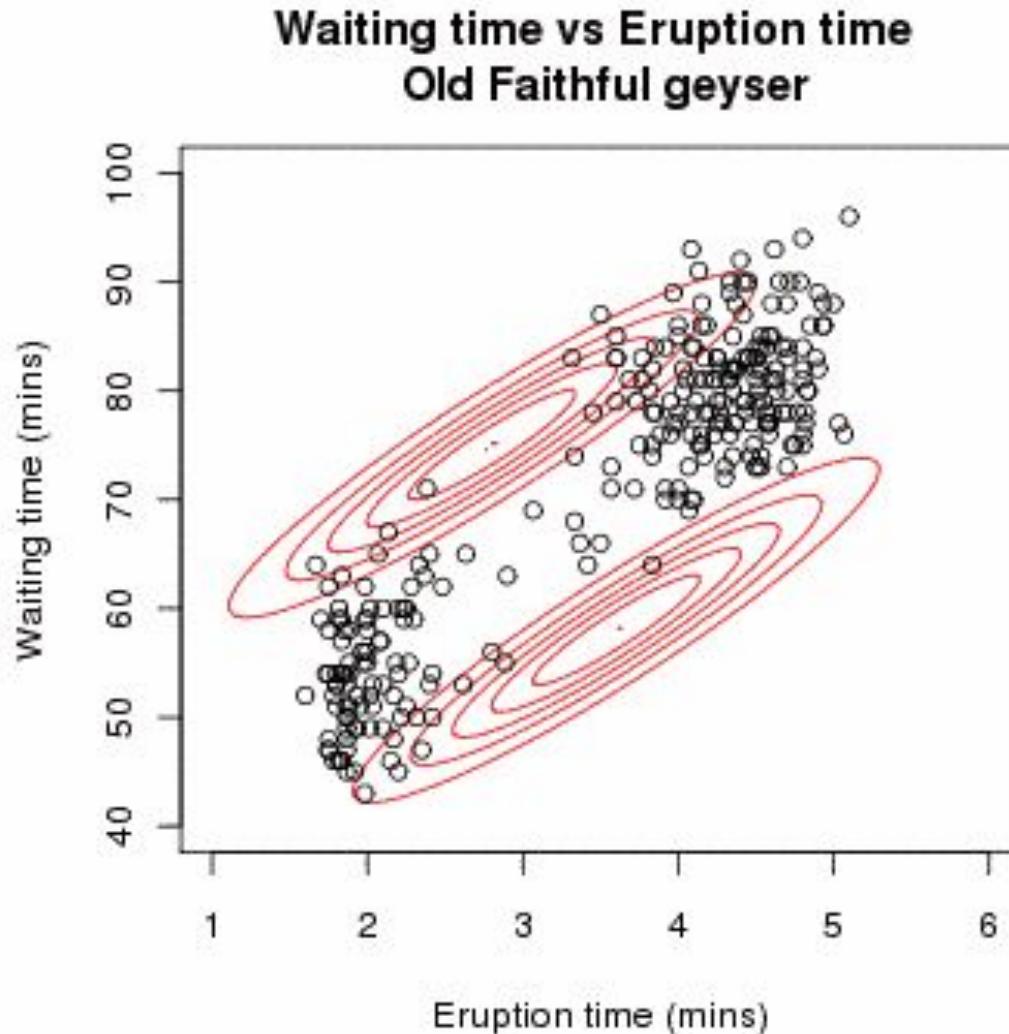


- Recursive procedure:
 1. Start: Given K , provide the initial parameters: $\{\pi_k, \mu_k, \Sigma_k\}$, $k=1\dots K$
 2. Classify: given $\{\pi_k, \mu_k, \Sigma_k\}$ compute the $\{\gamma_k\}$
 3. Re-center: Given $\{\gamma_k\}$, compute the $\{\pi_k, \mu_k, \Sigma_k\}$
 4. Repeat 2-3 until convergence
- This can be seen as Expectation Maximization (EM) framework!
Classification: **E**xpectation step
Re-centering: **M**aximization step (ML estimation)

Slide credit: Dr. Antonio M. López



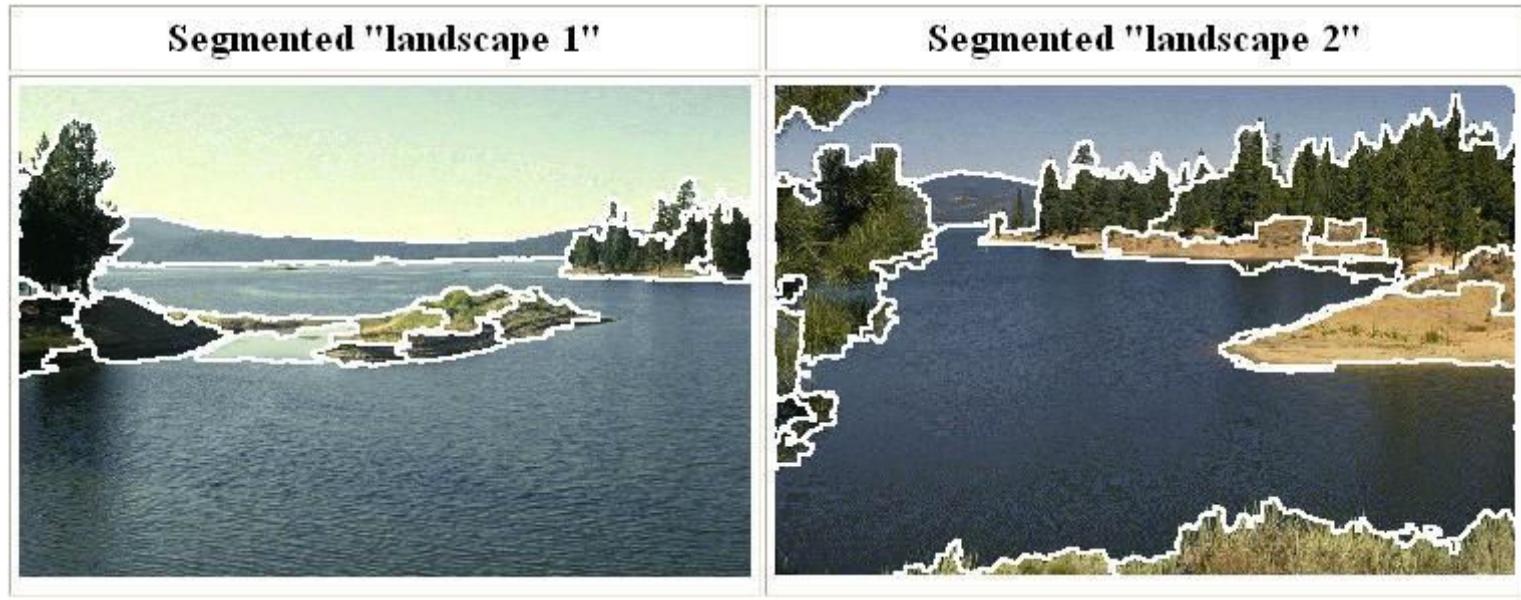
Homogeneity based – FS segmentation : GMM



http://en.wikipedia.org/wiki/Expectation_maximization

Homogeneity based – FS segmentation : Mean shift

- Technique for clustering-based segmentation
- It seeks nodes (local maxima of density) in feature space

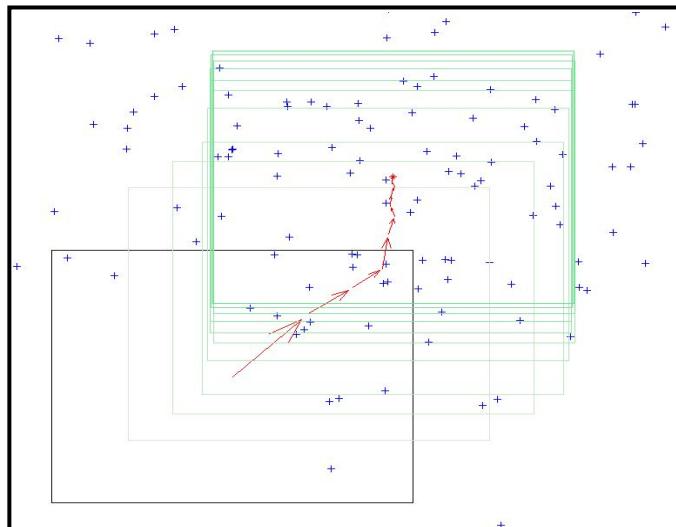


D. Comaniciu and P. Meer, Mean Shift: A Robust Approach toward Feature Space Analysis, PAMI 2002.

Homogeneity based – FS segmentation : Mean shift

Mean-shift

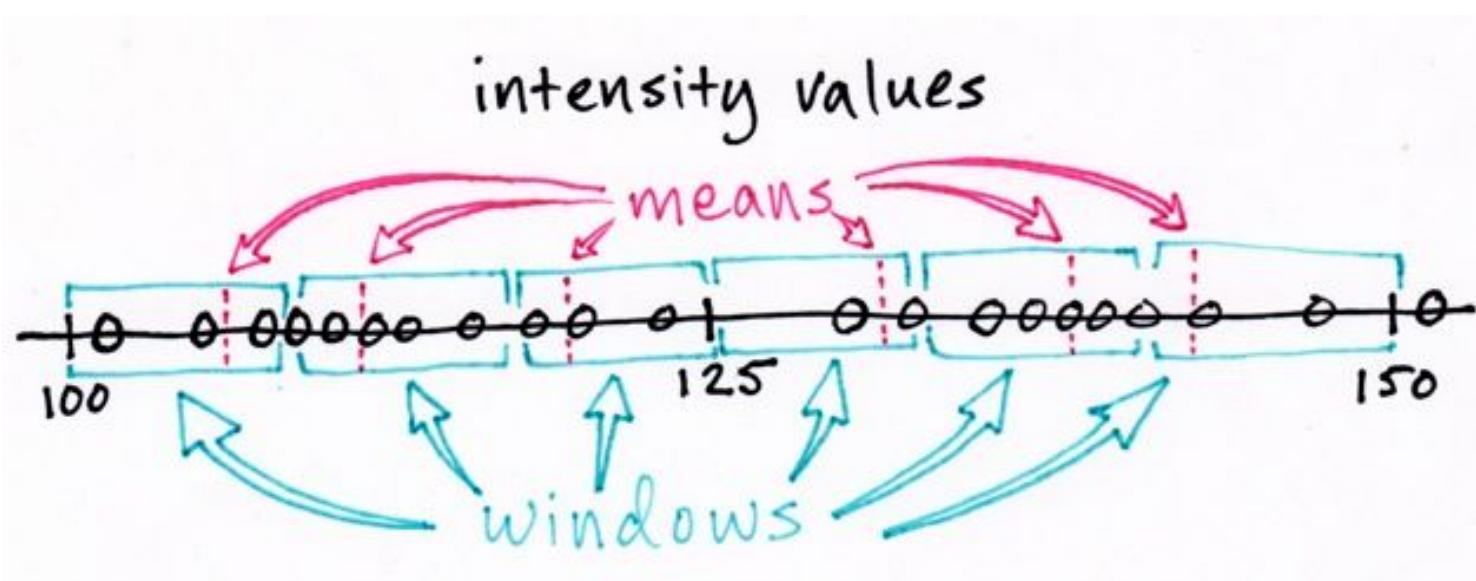
1. Select a search window size
2. Define a set of initial points for the search windows
3. Compute the mean location (centroid) of the data inside the window
4. Re-center the window at position computed in 3.
5. Iterate 3. and 4. until convergence



Slide credit: Gary Bradski & Sebastian Thrun

Homogeneity based – FS segmentation : Mean shift

1D example

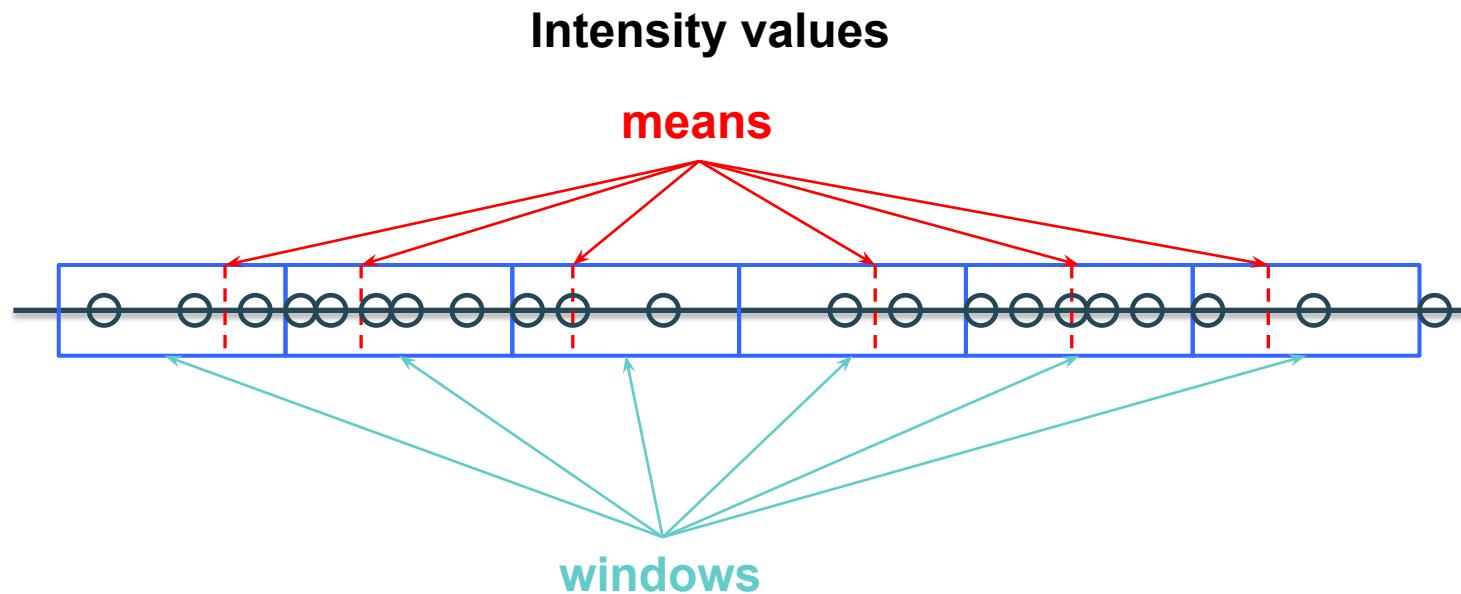


Source: Mike Laielli – benderseye.com



Homogeneity based – FS segmentation : Mean shift

1D example

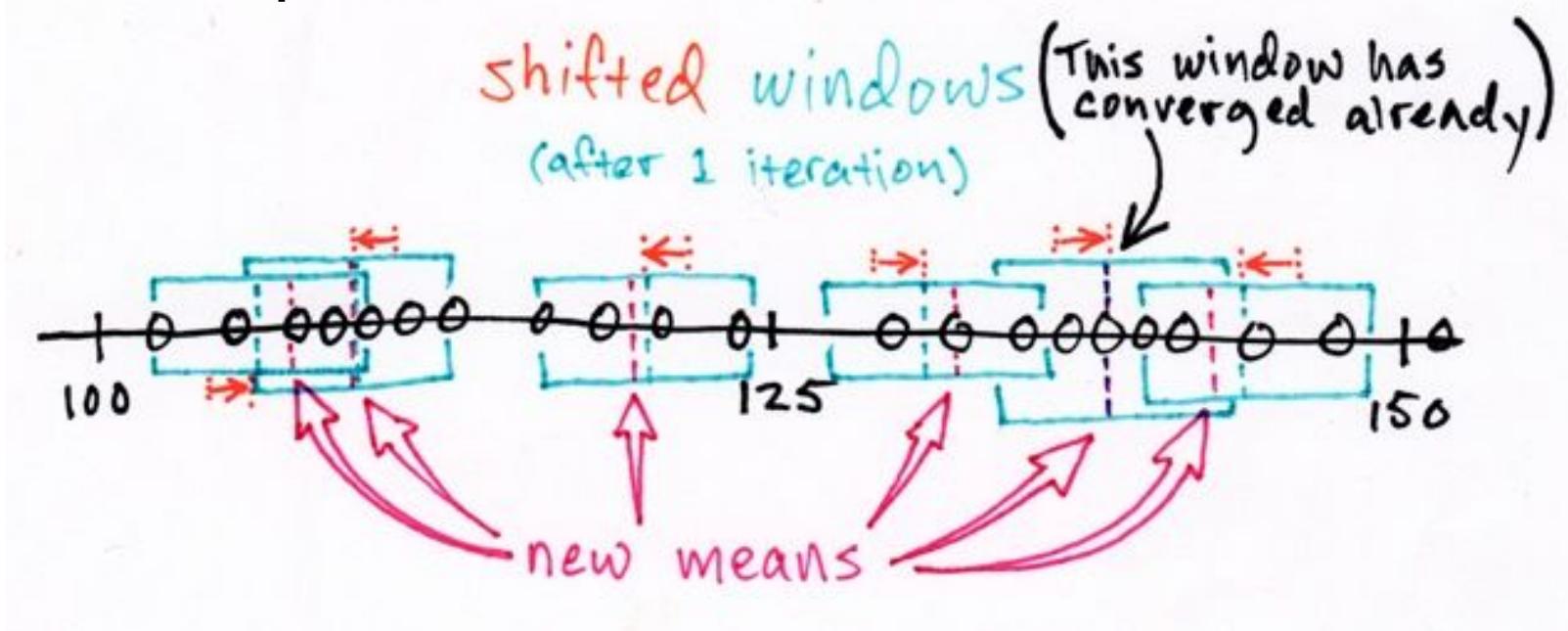


Source: Mike Laielli – benderseye.com



Homogeneity based – FS segmentation : Mean shift

1D example

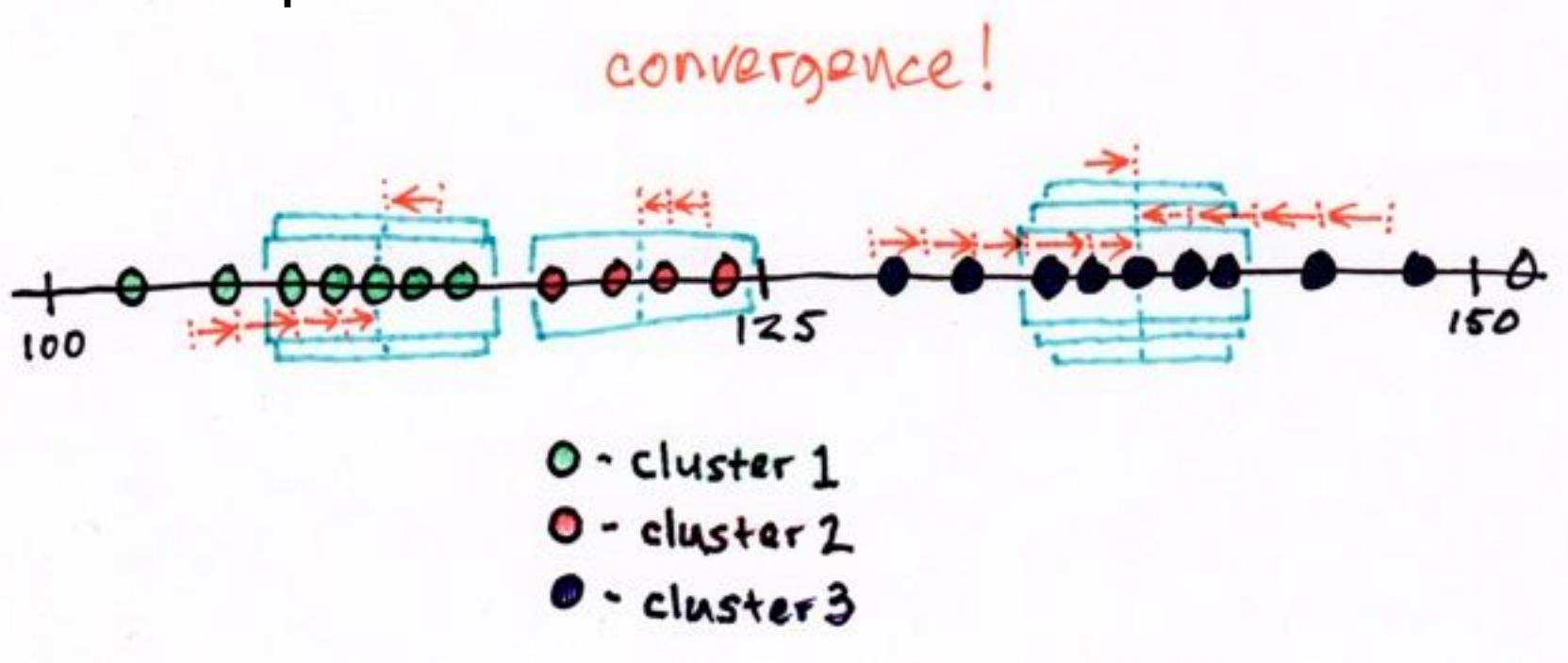


Source: Mike Laielli – benderseye.com



Homogeneity based – FS segmentation : Mean shift

1D example



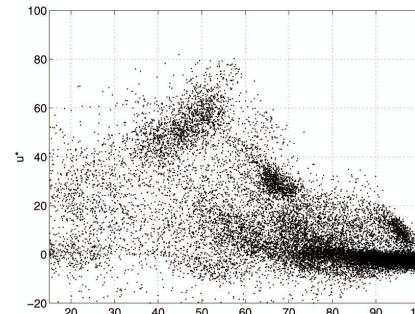
Source: Mike Laielli – benderseye.com



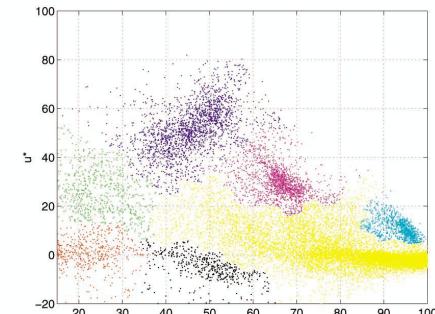
Homogeneity based – FS segmentation : Mean shift

Procedure:

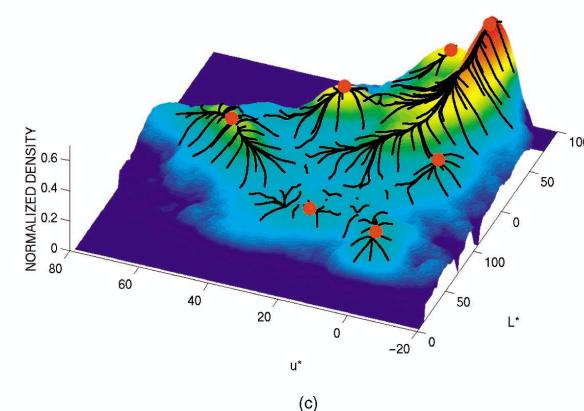
- Find features: (color, gradient, etc.)
- Select window size. Initialize windows at individual feature points
- Perform mean-shift for each window until convergence
- Merge windows that end near the same mode.



(a)



(b)



(c)

Homogeneity based – FS segmentation : Mean shift

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
(curse of dimensionality)

$$O(Tn^2)$$

T: Number of iterations
N : number of points in the dataset



Segmentation - FS segmentation : Conclusions

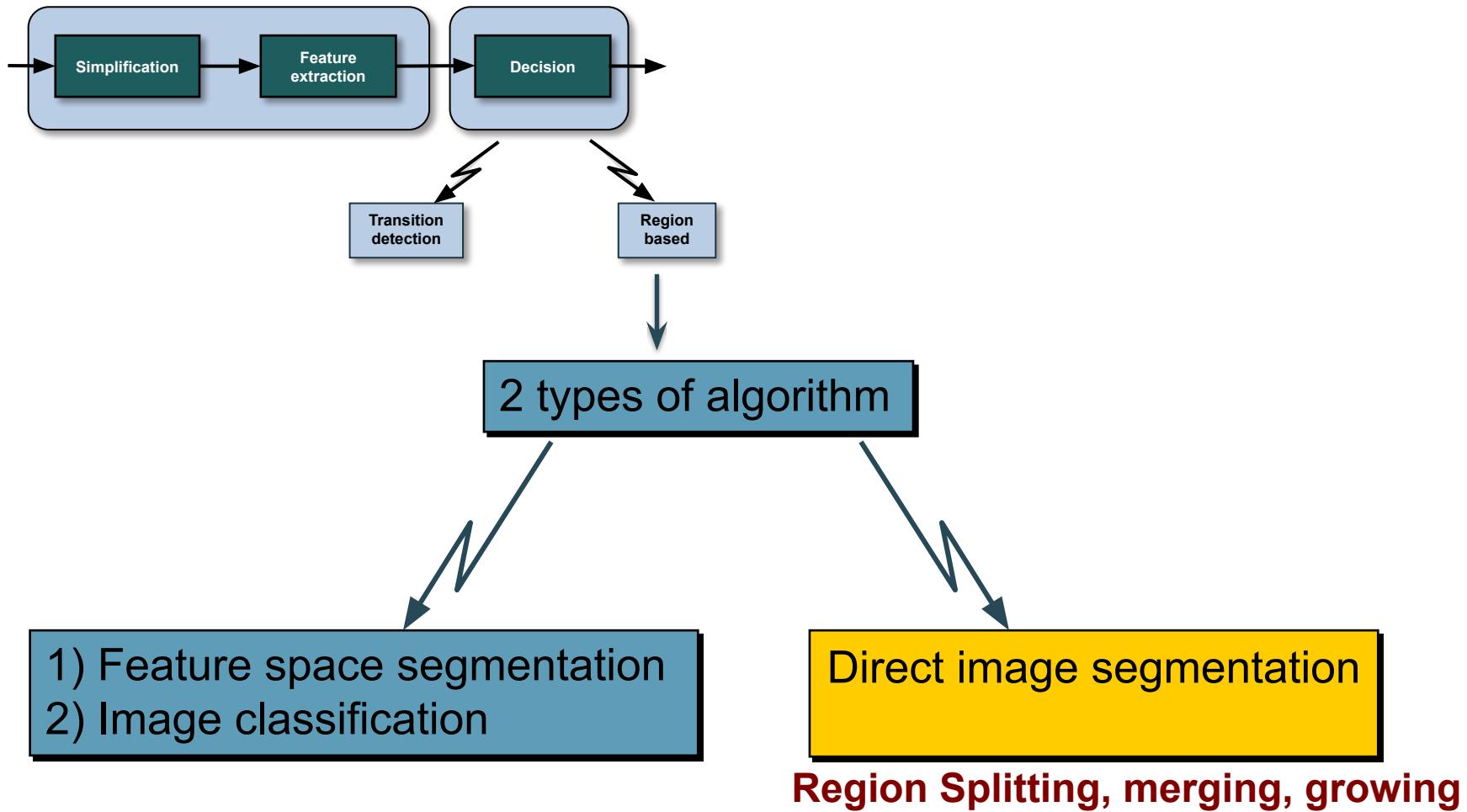
Strong point:

- Simple

Drawback:

- No control of the space connectivity
- May result on large number of **small isolated regions**

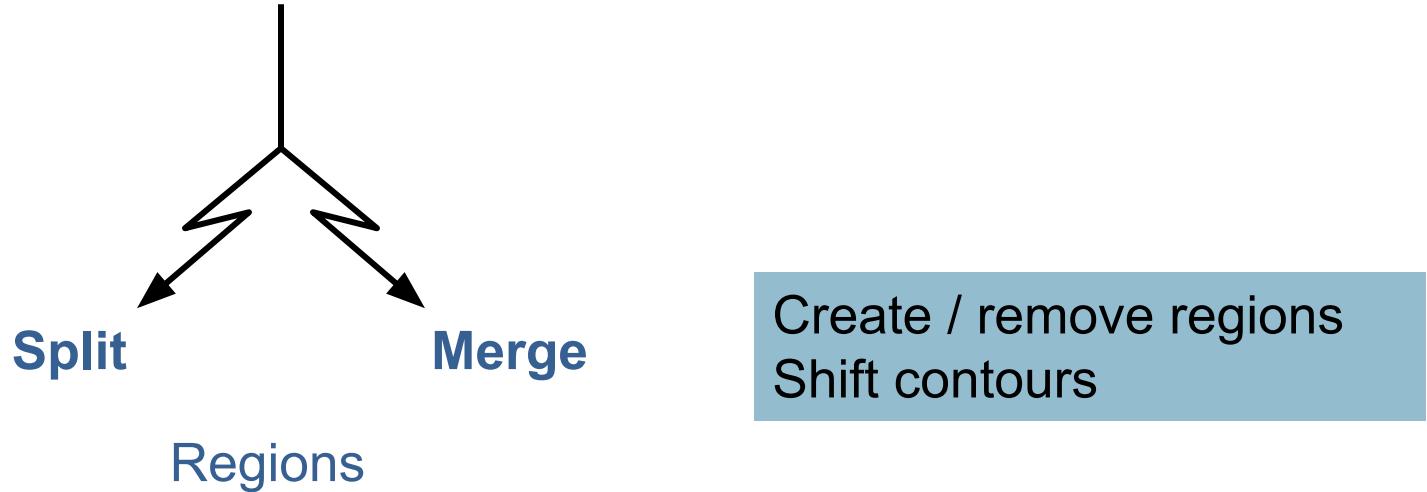
Segmentation: Homogeneity based



Homogeneity based – Region splitting, merging & growing

Strategy:

- Create an initial partition
- Define an optimization criterion
- Optimize (modify) the partition (criterion minimization)



Homogeneity based

Segmentation: Create a partition of the image such that

$$\begin{cases} C(R_i) = \text{True}, & \forall i \\ C(R_i \cup R_j) = \text{False}, & \forall i, j \end{cases}$$

Regions should group pixels which are homogenous.

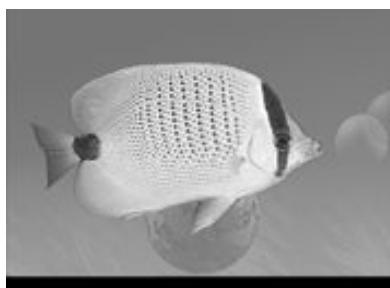
How to estimate the homogeneity?

- Define a region model
- Measure the variability of the pixels with respect to the model



Homogeneity based: region model

$x(i, j)$



R_n

$M_n(i, j)$



0 Order
Polynomial
(Mean)

$$M_n(i, j) = \sum_{i, j \in R_n} x(i, j)$$



1st order
polynomial

$$M_n(i, j) = ai + bj + c$$



2nd order
polynomial

$$M_n(i, j) = ai^2 + bj^2 + cij + di + ej + f$$

Homogeneity based: measure of variability

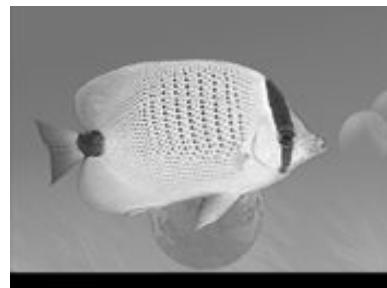
Classical example: Mean Squared Error

$$C_T = \frac{1}{N} \sum_n \sum_{i,j \in R_n} [x(i,j) - M_n(i,j)]^2$$

Gray level
(or color)

- Region model:
- Constant (mean)
 - Polynomial, ...

R_n : Region
i,j: Position
N: Pixel num.



0 Order
polynomial

But is this enough?

Segmentation: Homogeneity based

4 Examples:

- Split & Merge (top-down)
- Region merging (bottom-up)
- Region growing (seeds)
 - Watershed

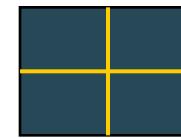


Homogeneity based – Split & Merge

- || 1) Initial partition:
 - || 2) Classical criterion, C:
 - || Image = 1 region (top-down)
|| Mean squared error with respect
|| to a 0, 1st or 2nd order model
-

“Split”:

If $C(R_i) > T_0$ → Geometric splitting

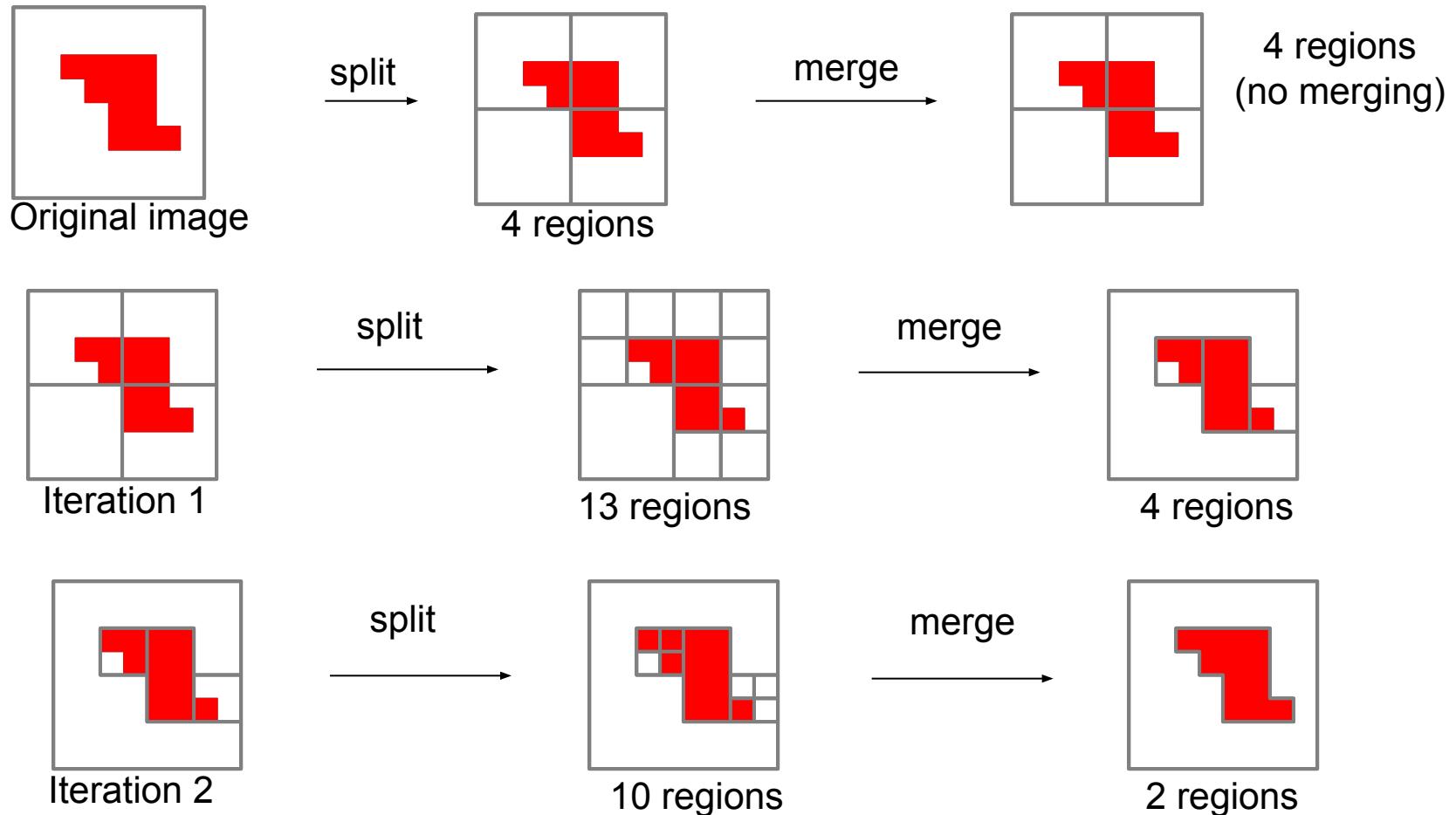


“Merge”:

If $C(R_i \cup R_j) < T_1$ → Merge R_i and R_j



Homogeneity based – Split & Merge

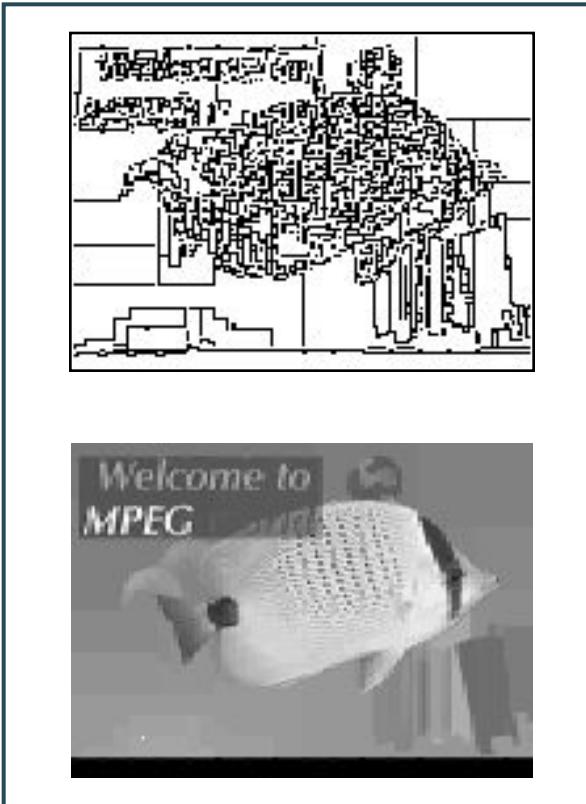


Split → If region is not homogeneous, split in 4

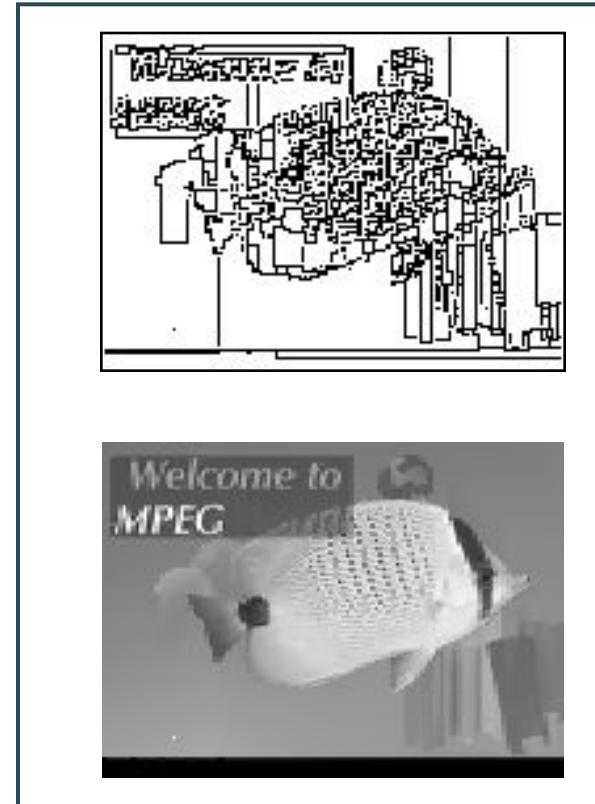
Merge → If neighboring regions are homogenous, Merge them

Homogeneity based – Split & Merge

0 order model



2nd order model



Split & Merge:

- Simple
- Simple initial partition
- Global view

- Pure geometrical split
- Contour are not always natural

Segmentation: Homogeneity based

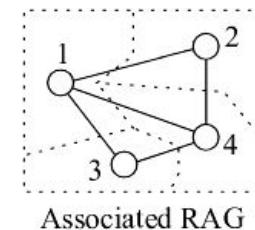
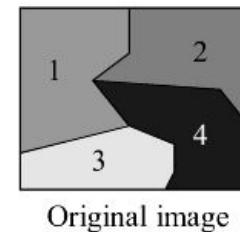
4 Examples:

- Split & Merge (top-down)
- Region merging (bottom-up)
- Region growing (seeds)
 - Watershed

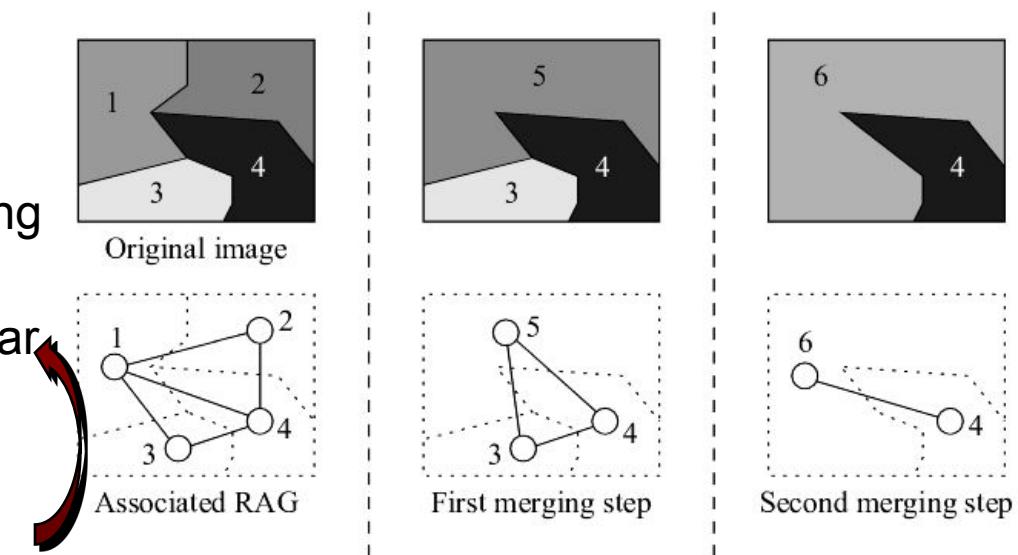


Homogeneity based – Region merging

- Region Adjacency Graph (RAG)
 - Node: regions of the image
 - Edge: Neighbor relationship



- Iterative merging process:
 - Define an initial partition and its RAG (nodes: regions, edges: neighboring region similarity)
 - Find the pair of most similar neighboring regions
 - Merge them
 - Update the edges



Homogeneity based – Region merging

- Initial partition:
 - Each pixel is an individual region
 - Over-segmentation
- Similarity between regions:
 - Approximation of Mean Squared Error:

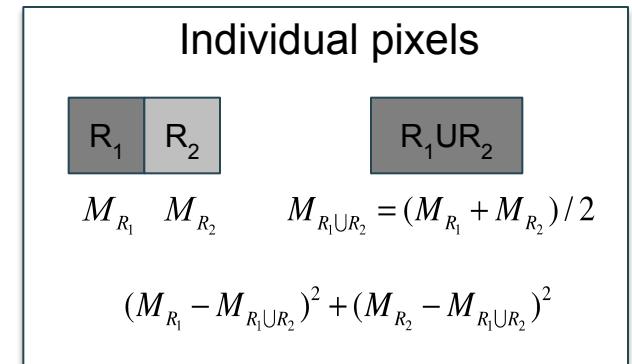
$$C_{color}(R_1, R_2) = N_{R1} \left\| M_{R1} - M_{R1 \cup R2} \right\|_2^2 + N_{R2} \left\| M_{R2} - M_{R1 \cup R2} \right\|_2^2$$

- Contour length variation:

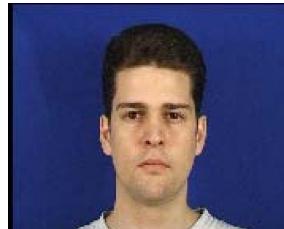
$$C(R_1, R_2) = \alpha C_{color}(R_1, R_2) + (1 - \alpha) C_{cont}(R_1, R_2)$$

$$C_{cont}(R_1, R_2) \approx -\text{Length of common contour}$$

- Stopping criterion:
 - Given number of regions
 - PSNR resulting from modeling the regions of the partition



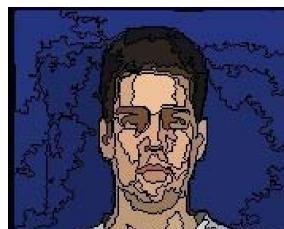
Homogeneity based – Region merging



Region
number (50)



22,00 dB [50 regions]



29,50 dB [50 regions]



22,13 dB [50 regions]



32,29 dB [50 regions]

PSNR (26dB)



[26 dB] 273 regions



[26 dB] 13 regions



[26 dB] 347 regions



[26 dB] 7 regions

Slide credit: Verónica Vilaplana

Segmentation: Homogeneity based

4 Examples:

- Split & Merge (top-down)
- Region merging (bottom-up)
- Region growing (seeds)
 - Watershed

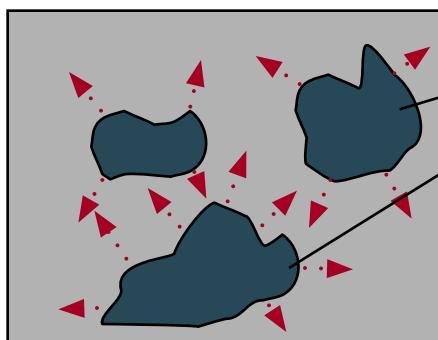
Homogeneity based – Region growing

Basic strategy:

- Initial segmentation: Partial segmentation

$$\bigcup_i R_i \neq E$$
$$E \setminus \bigcup_i R_i = \text{Uncertainty zone}$$

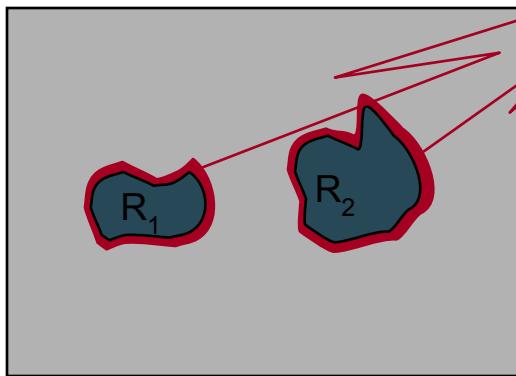
- Progressive elimination of the uncertainty zone



Initial regions:

- Regions interior
- “Safe” areas
- Application dependent

Homogeneity based – Region growing



Pixels from the uncertainty area that are connected to regions: $\{p_j^{R_i}\}$

Assign iteratively the “closest” pixel

$p_j^{R_i}$ is assigned to R_i if:

$$C\{R_1, \dots, \{p_j^{R_i} \cup R_i\}, \dots, R_k\} < C\{R_1, \dots, \{p_m^{R_n} \cup R_n\}, \dots, R_k\}$$

$$\forall (n, m) \neq (i, j)$$

⇒ Merging between regions and pixels from the uncertainty area

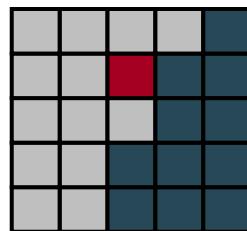
Homogeneity based – Region growing

Criterion: Local estimation of the variation of the criterion

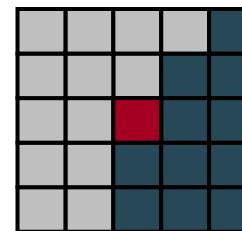
For each potential merging between p_0 and R_i , compute:

$$\Delta C = \alpha (p_0 - M_{R_i})^2 + (1-\alpha) \Delta \text{contour}$$

Pixel gray level Region mean Variation of the contour length

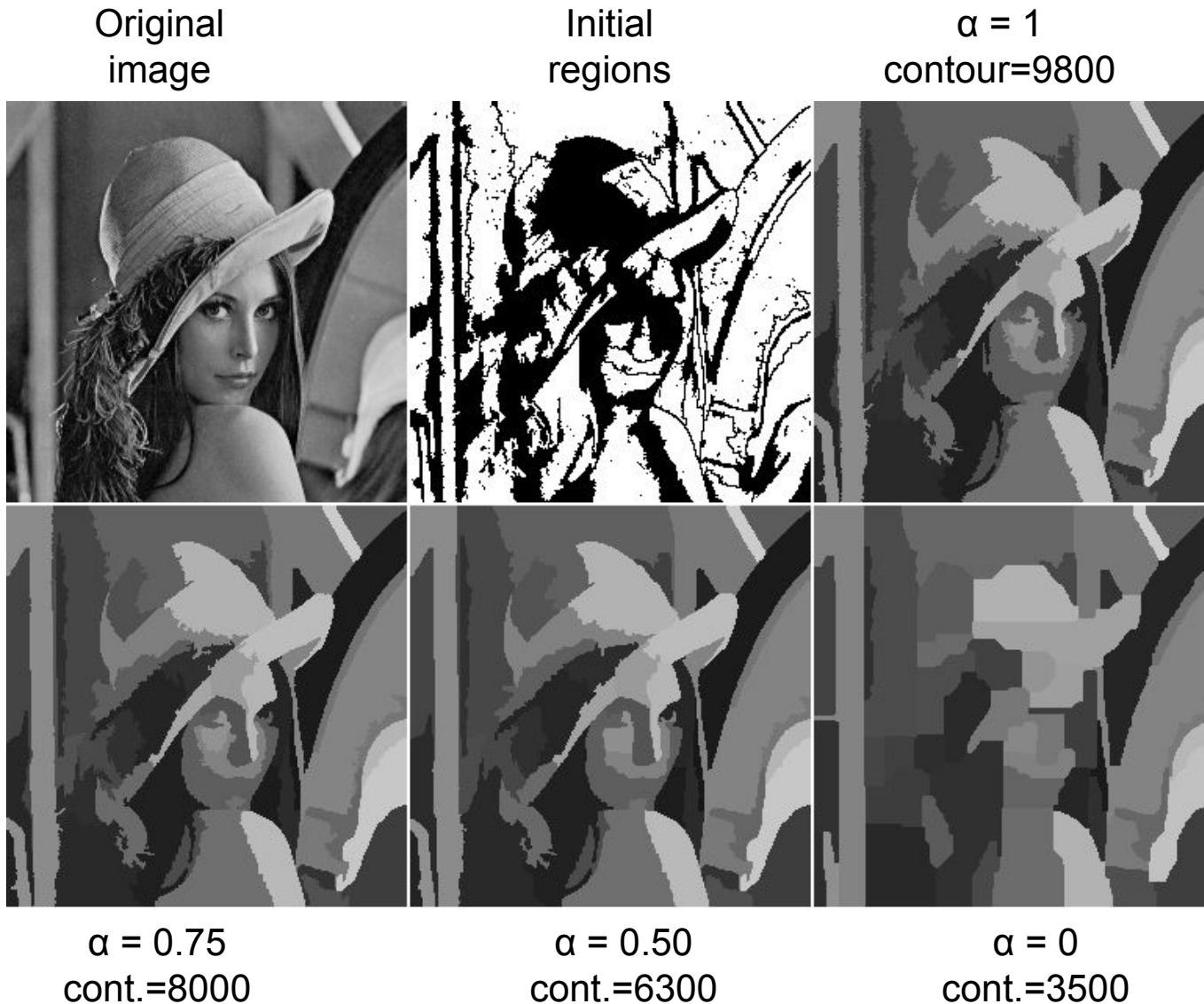


$$\Delta \text{contour} = 3-1 = 2$$



$$\Delta \text{contour} = 2-2 = 0$$

Homogeneity based – Region growing

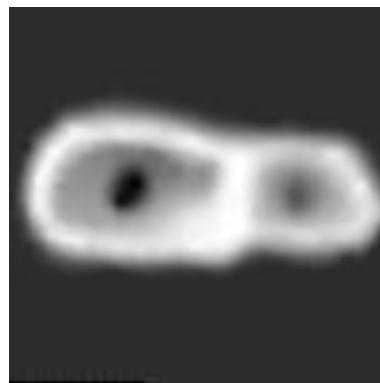


Homogeneity based – Region growing : Watershed

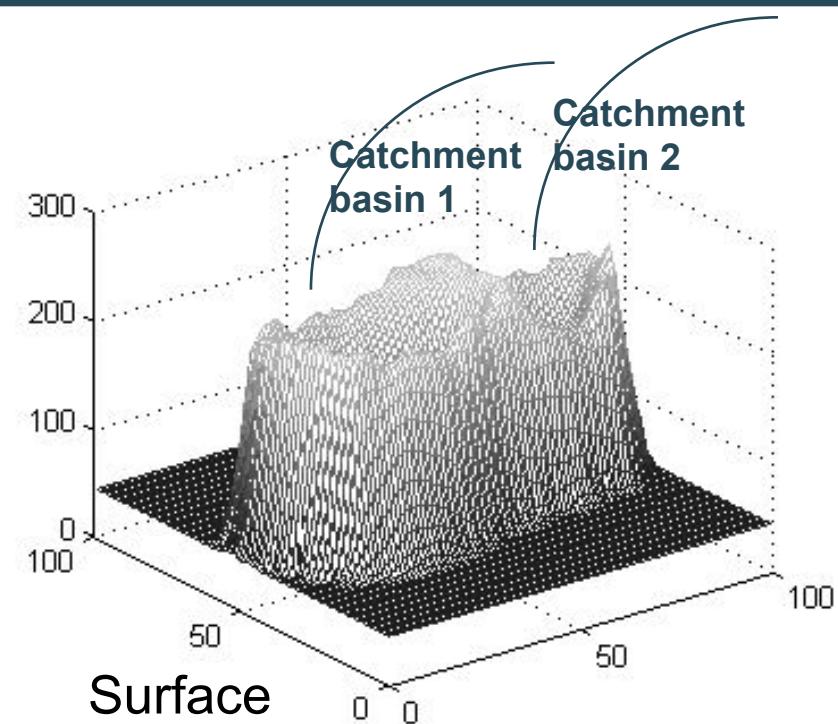
Watershed: Morphological approach to image segmentation through region growing

Image → Topographic surface

Gray level → Height

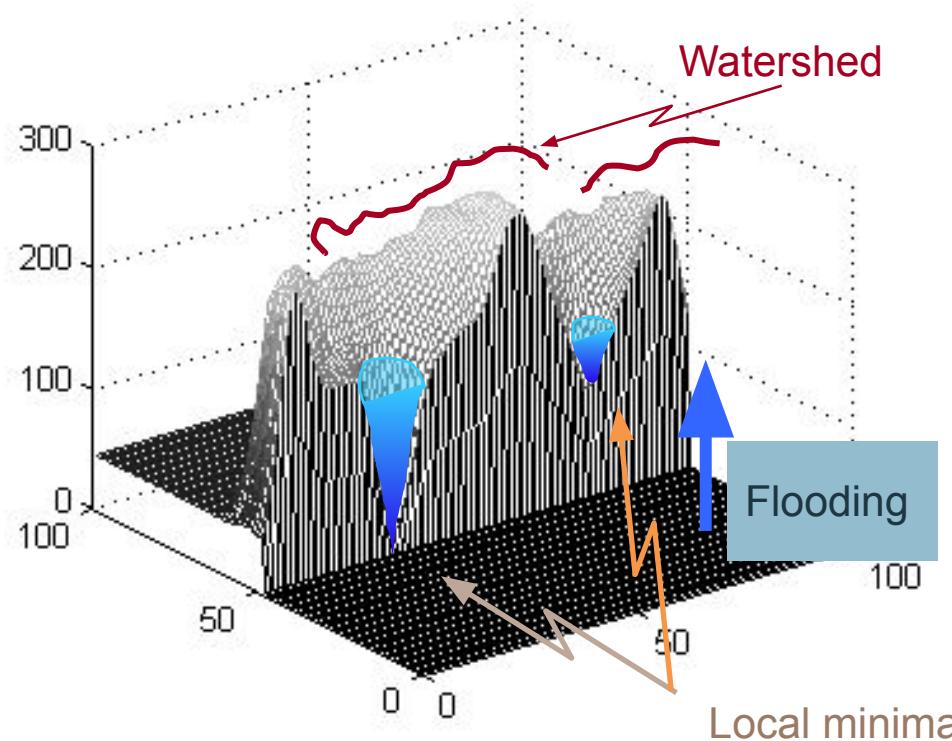


Original
image



Homogeneity based – Region growing : Watershed

Watershed: locations where 2 (or more) catchment basins meet



Each pixel is assigned to the catchment basin assigned to its minimum
→ Flooding algorithm

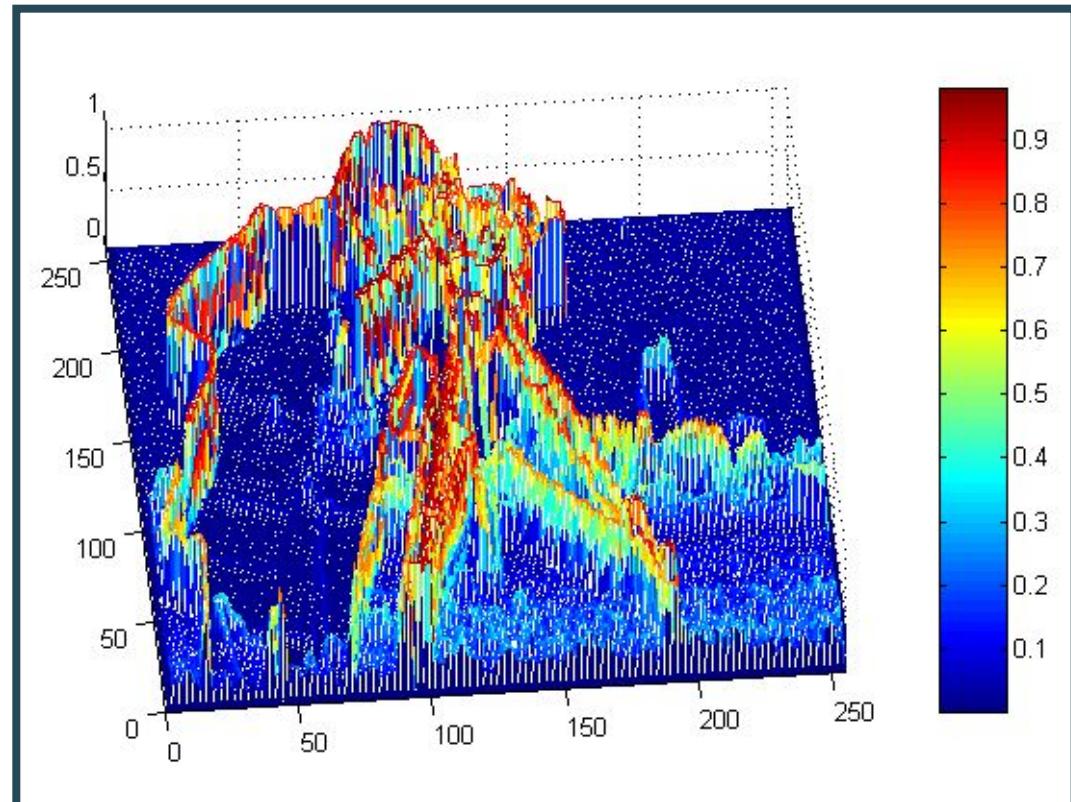
Homogeneity based – Region growing : Watershed



Original image



Gradient



→ image contours = watershed of gradient

Homogeneity based – Region growing : Watershed

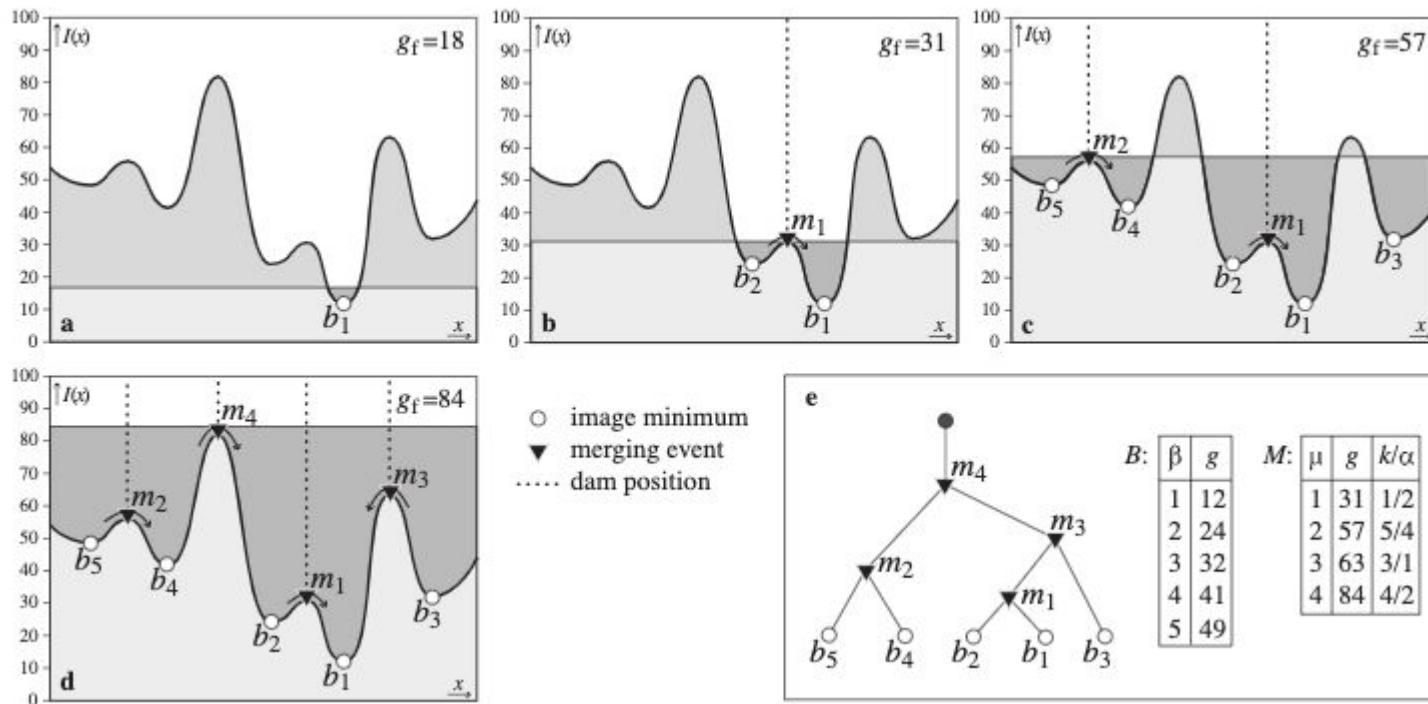
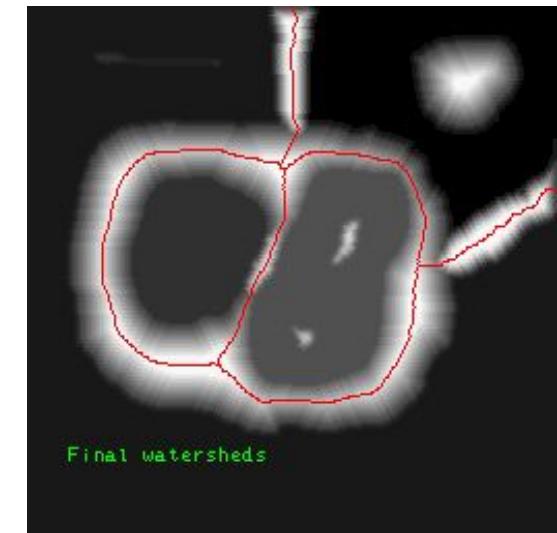
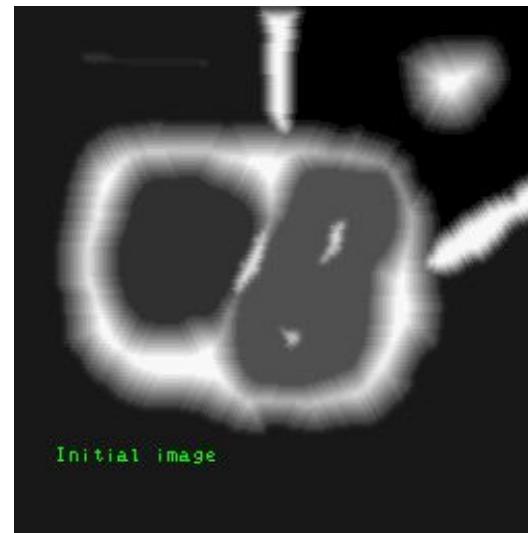
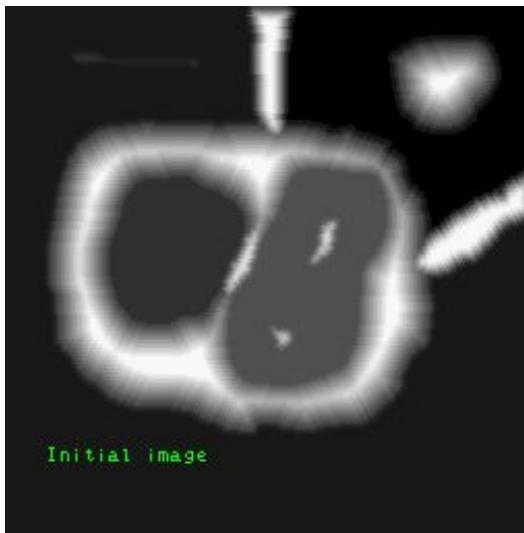


Illustration of hierarchical watershed transform on a continuous 1d function $I(x)$ interpreted as a landscape. The landscape is sequentially flooded from bottom to top. **a:** At a flooding height $g_f = 18$, a single basin b_1 exists. **b:** At $g_f = 31$, a second isolated minimum b_2 and the lowest separating ridge have been detected. At the ridge point, a candidate for basin merging m_1 is registered and stored for future evaluation. **c:** At higher flooding heights, more and more image minima and merging procedures are registered. **d:** At $g_f = 84$ the flooding is complete. **e:** All (atomic) basins and merging candidates have now been detected and are ordered in a hierarchical tree. Relevant information, i.e., the respective gray levels $g[\beta]$ and $g[\mu]$ and merging candidates $k[\mu]$ and $\alpha[\mu]$, are stored in the basin and merging tables B and M

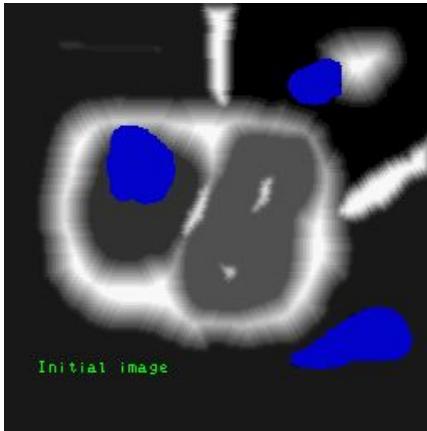
Horst K. Hahn and Heinz-Otto Peitgen, "IWT – Interactive Watershed Transform: A hierarchical method for efficient interactive and automated segmentation of multidimensional grayscale images", Proc. Medical Imaging, SPIE 5032, San Diego, Feb 2003

Homogeneity based – Region growing : Watershed

- Initial regions: Local minima of the image
- Growing process: The pixel with the gray level that is the closest to the flooded level is assigned to its corresponding catchment basin.



Homogeneity based – Region growing : Watershed



Original image with marker



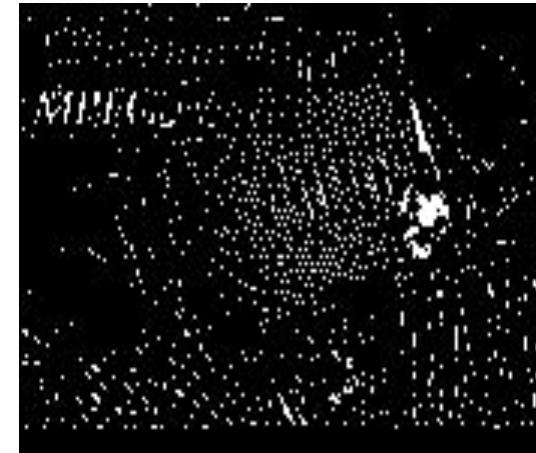
Markers: initial partition



Homogeneity based – Region growing : Watershed



Origina



Maxim



Homogeneity based – Region growing

Advantages:

- Precise contours
- Control the region connectivity

Drawbacks:

- Need of a partial segmentation
 - Markers
 - Definition of a “safe” area

Mixed transition/homogeneity – gPb-OWT-UCM

- Ultrametric contour maps (UCM)
 - Transforms a contour map into a hierarchy of regions
 - Iterative merging using *Ultrametric* dissimilarities



Mixed transition/homogeneity – gPb-OWT-UCM

Original image

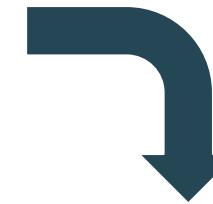
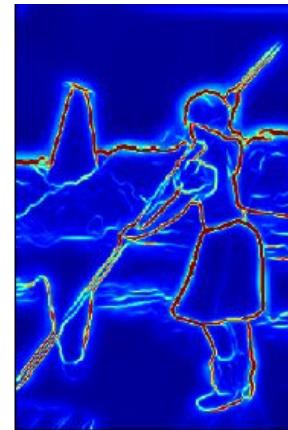


- Local cues
- Global cues



gPb contour
probabilities

Contour image

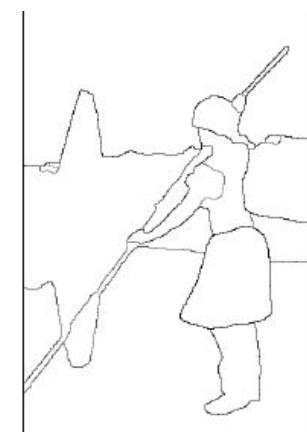


Oriented Watershed
Transform

gPb: D. Martin, C. Fowlkes, and J. Malik, "Learning to detect natural image boundaries using local brightness, color and texture cues," PAMI, 2004.



Iterative
merging

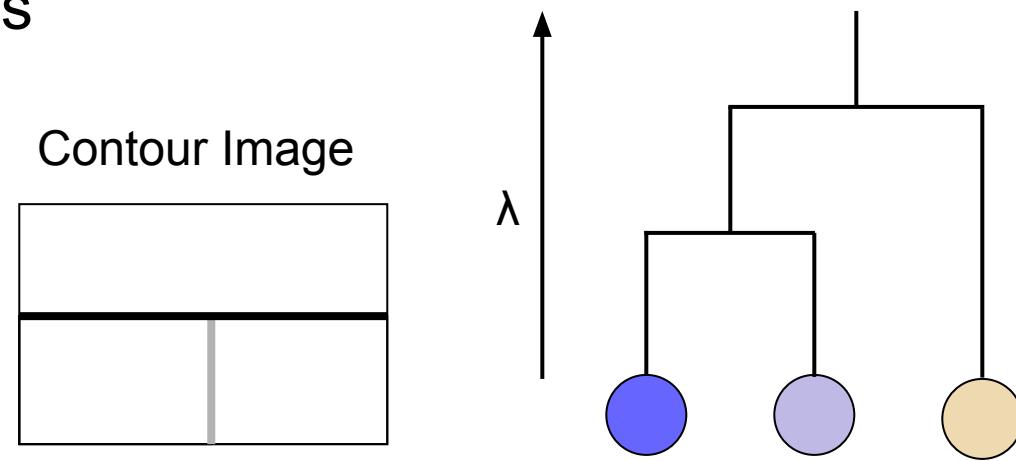


P. Arbelaez, M. Maire, C. Fowlkes, J. Malik. **Contour Detection and Hierarchical image Segmentation**. IEEE Trans. on PAMI , 2011.

Source: Hsin-Min Cheng

Mixed transition/homogeneity – gPb-OWT-UCM

- Defines a duality between closed, non-self-intersecting weighted contours and a hierarchy of regions
- The base level of this hierarchy respects even weak contours and is thus an over-segmentation of the image.
- Upper levels of the hierarchy respect only strong contours, resulting in an under-segmentation.
- Moving between levels offers a continuous trade-off between these extremes



Mixed transition/homogeneity – gPb-OWT-UCM

- Define an initial graph $G = (P_0; K_0; W(K_0))$
 - The nodes are the regions, P_0
 - The links are the arcs K_0 separating adjacent regions
 - The weights $W(K_0)$ are a measure of dissimilarity between regions
- Dissimilarity between two adjacent regions is defined as the average strength of their common boundary in K_0

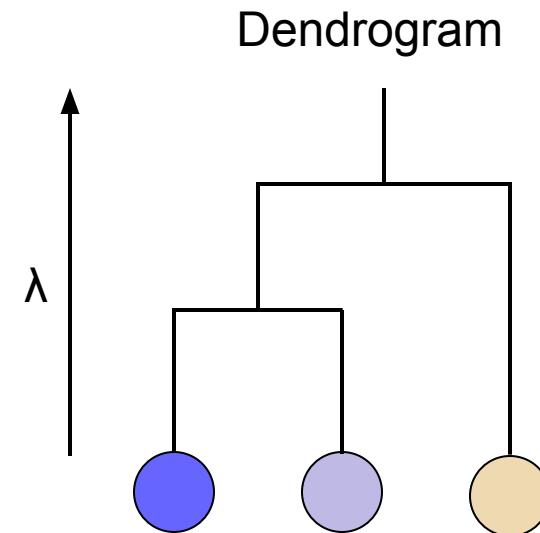
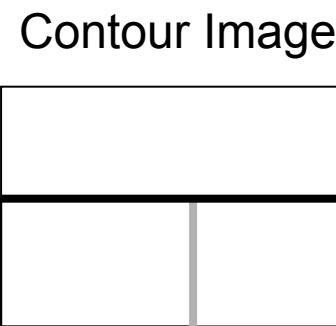
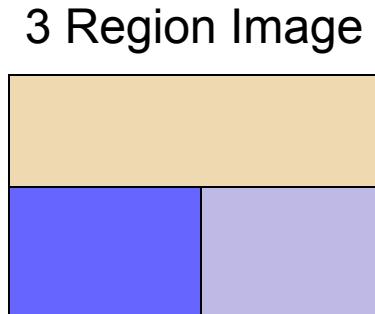
Algorithm:

- 1) Select minimum weight contour:
 $C^* = \operatorname{argmin}_{C \in \mathcal{K}_0} W(C).$
- 2) Let $R_1, R_2 \in \mathcal{P}_0$ be the regions separated by C^* .
- 3) Set $R = R_1 \cup R_2$, and update:
 $\mathcal{P}_0 \leftarrow \mathcal{P}_0 \setminus \{R_1, R_2\} \cup \{R\}$ and $\mathcal{K}_0 \leftarrow \mathcal{K}_0 \setminus \{C^*\}.$
- 4) Stop if \mathcal{K}_0 is empty.
Otherwise, update weights $W(\mathcal{K}_0)$ and repeat.



Mixed transition/homogeneity – gPb-OWT-UCM

- Hierarchical segmentation



Source: Pablo Arbeláez

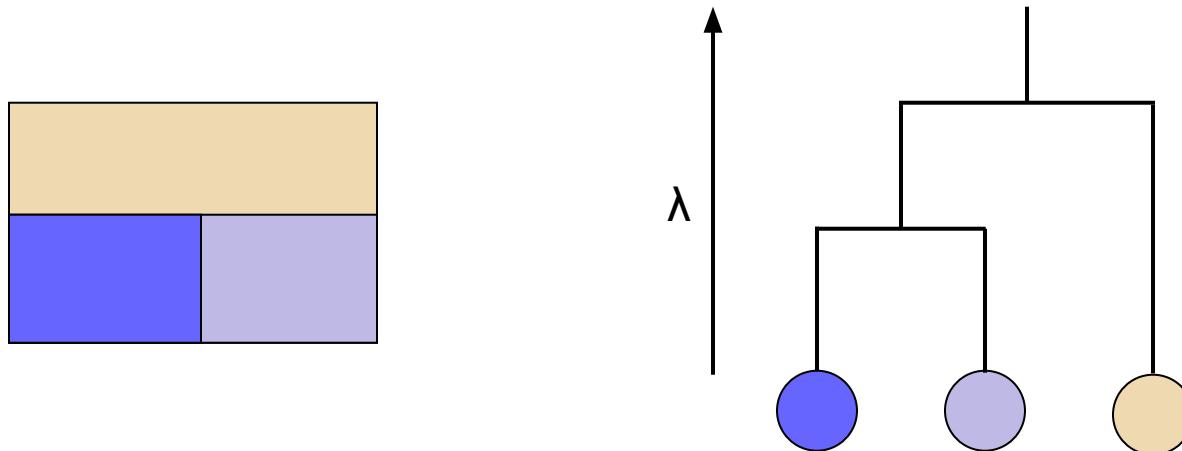


Mixed transition/homogeneity – gPb-OWT-UCM

- Ultrametric

$$D(R_1, R_2) \leq \max \{D(R_1, R), D(R, R_2)\}$$

The union R_{12} of two regions R_1 and R_2 must have \geq distance to adjacent region R than either R_1 or R_2

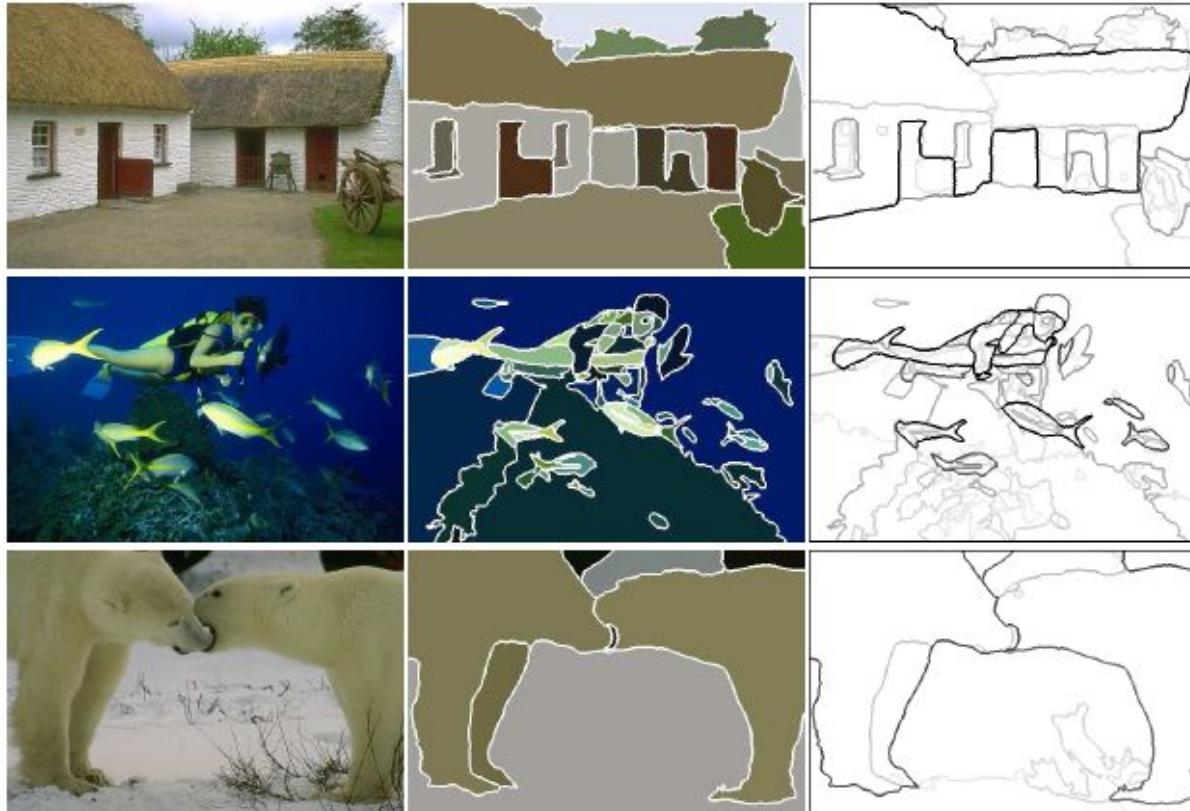


Source: Pablo Arbeláez



Mixed transition/homogeneity – gPb-OWT-UCM

Results



Source: Pablo Arbeláez

Fast UCM implementation (MCG Pre-trained, Matlab/C++):

<http://www.eecs.berkeley.edu/Research/Projects/CS/vision/grouping/mcg/>

Segmentation: 2019 state of the art

- Research focused on semantic segmentation, instance segmentation



- Mostly deep learning

Semantic segmentation: popular architectures:

<https://towardsdatascience.com/semantic-segmentation-popular-architectures-dff0a75f39d0>

A 2019 Guide to Semantic Segmentation:

<https://heartbeat.fritz.ai/a-2019-guide-to-semantic-segmentation-ca8242f5a7fc>

- Classical methods still used:

- Max-tree representation

Yongchao Xu, Edwin Carlinet, Thierry Géraud, Laurent Najman. Hierarchical Segmentation Using Tree-Based Shape Spaces. IEEE Transactions on PAMI, 2017, 39 (3)

Salembier P, Liesegang S, López-Martínez C. Ship Detection in SAR Images Based on Maxtree Representation and Graph Signal Processing. IEEE Transactions on Geoscience and Remote Sensing. 2019;57(5)

Segmentation: Summary

