Image Segmentation

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Reference Books:

• Computer Vision: Algorithms and Applications. Richard Szeliski. Springer. 2010.

http://szeliski.org/Book

Content

- Introduction
- Contour-based techniques
- Thresholding
- Region-based techniques
 - Region growing
 - K-Means
 - Expectation-Maximixation
 - Mean-Shift (not included)
- Semantic segmentation

1. Introduction

Segmentation ...

• divides an image in regions whose pixels have similar properties (intensity, color, texture, location in the image, ...).

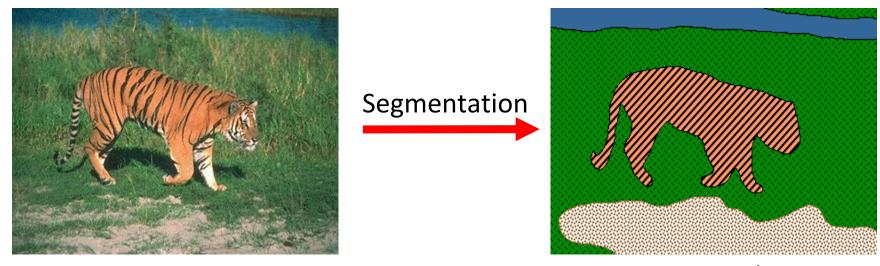


Image segmented in 4 regions

1. Introduction

Conceptually, two approaches exist:

■ Top down (Semantic segmentation): Pixels from the same scene object should be in the same region



The zebras are one object!

■ Bottom up (pixel-driven segmentation): Similar pixels must be in the same region



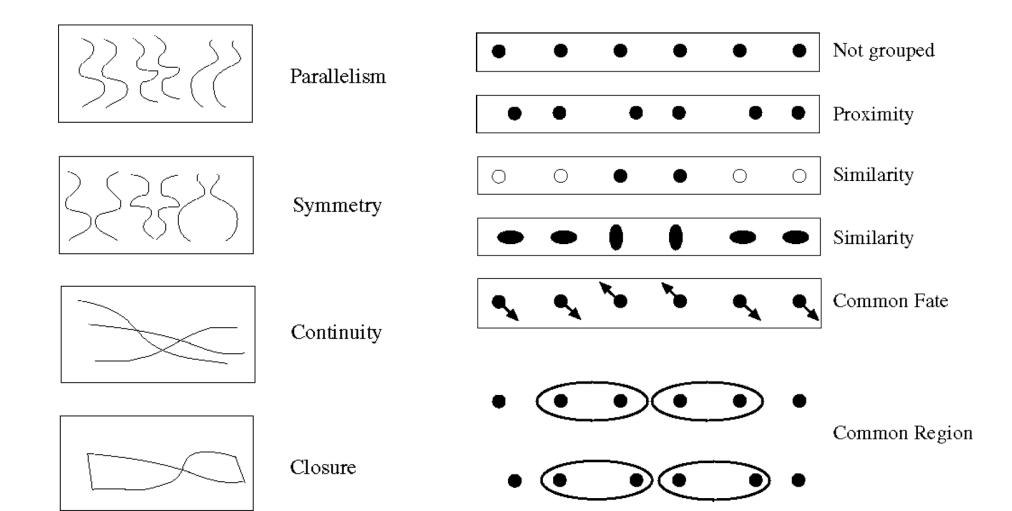
Each black/white band is an object!

Different results depending on the properties/knowledge employed



1. Introduction

The Gestalt theory stablishes properties/rules that humans employ to group entities of the images:



Examples of human segmentation

Using similarity, proximity



Using knowledge of the world



If we don't know about the features of a dalmatian dog, no way to segment this image!

2. Contour-based techniques

Atempt to indentify the image regions by detecting the their contours

Image contours: edge pixels that enclose a region of similar intensities

Two main approaches:

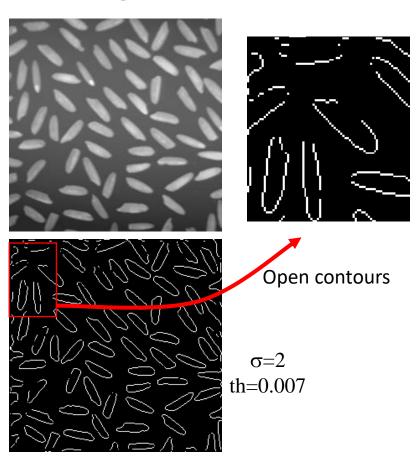
Local techniques → { LoG + zero crossing Edge following → Canny operator

Global techniques

Hough transform

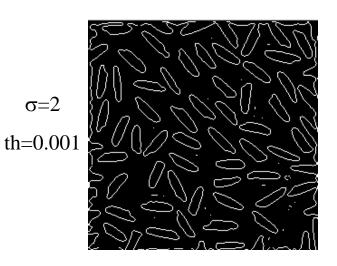
2. Contour-based techniques

LoG + Zero crossing



MATLAB

```
I = imread('rice.tif');
%BW=edge(I,'log',thresh,sigma)
log = edge(I,'log',0.001,2);
figure, imshow(log)
```

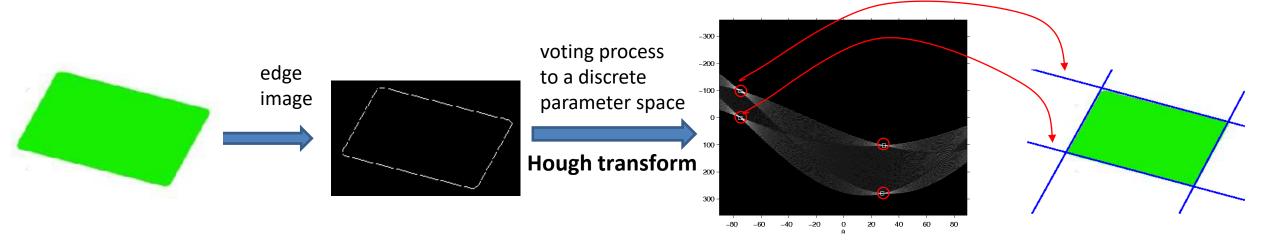


It's an interesting method for region segmentation since closed contours provide regions of **similar pixel intensities**.

2. Contour-based techniques

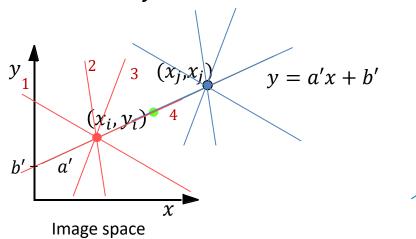
Hough transform

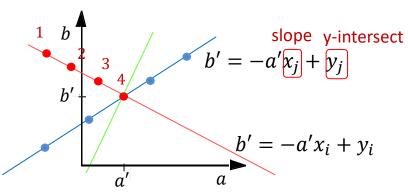
- Algorithm patented by Paul Hough to recognize lines in photographs (Hough, 1962)
- Can detect any shape in the image
 - Analytical forms (classic Hough): typically line, circle, ellipse.
 - Numerically described forms (generalized Hough): shape given by a table
- Based on a voting scheme: each point (x,y) of an edge image votes for a parameter set that defines the shape in the parameter space



Hough transform: Line detection

Basic form: y = ax + b



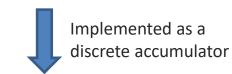


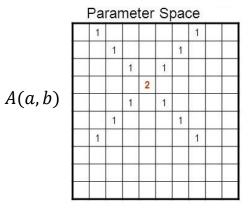
Parameter space: a, b

• Each point (x_i, y_i) transforms to a line in the (a, b)-parameter space

$$(x_i, y_i) \rightarrow b = -ax_i + y_i$$

• Lines through points (x_i, y_i) transform to points in the (a, b)-parameter space





Algorithm

- Quantize parameter space (a, b)
- Create Accumulator array A(a, b)
- Set $A(a,b) = 0 \ \forall a,b$
- For each (x_i, y_i) in the edge image For $a_k = 1$ to N $b_k = -a_k x_k + y_k$

$$b_k = -a_k x_i + y_i$$

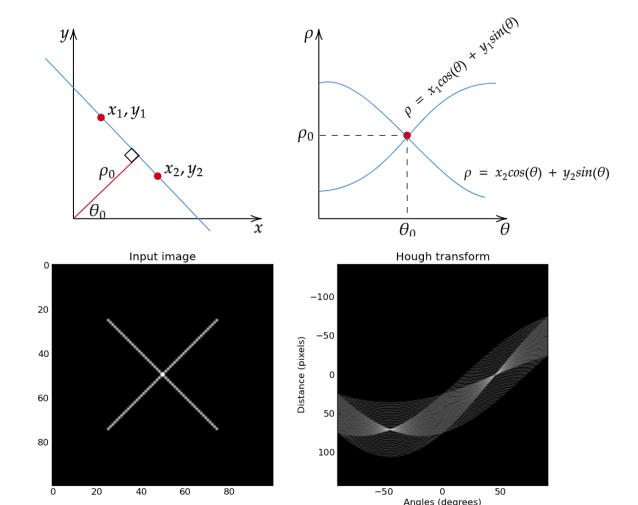
$$A(a_k, b_k) = A(a_k, b_k) + 1$$

• Find local máxima in A(a, b)

Hough transform: Line detection

Normal form: $\rho = x\cos\theta + y\sin\theta$

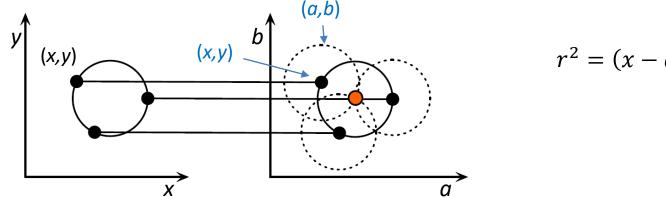
This line description does not have problems with vertical lines ($a = \infty$)



Hough transform

Example: Detecting circles

A circle of known radius r has two parameters: the coordinates of the center (a,b)

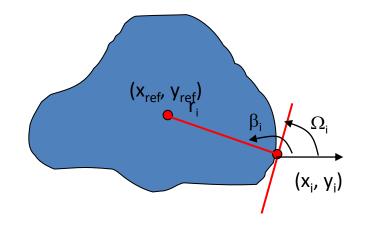


$$r^2 = (x - a)^2 + (y - b)^2$$

- Each point (x,y) in image space (left) votes for a circle (dashed) of candidate centers (a,b) in the parameter space (right).
- Ideally, all the generated circles in parameter space intersect at a certain (a,b) that is the unknown center of the circle in the image.

Hough transform

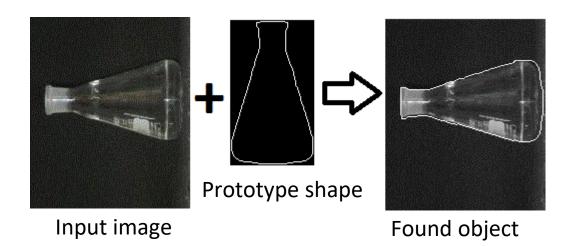
For numerically described shapes (generalized Hough): shape given by a table



Ω_1	
Ω_2	
•••	
Ω_{i}	$(r_1, \beta_1), (r_2, \beta_2)$

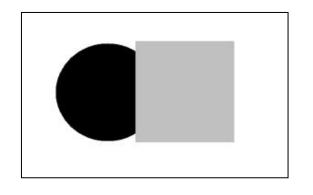
- For the tangent angle Ω_i at each contour point (x_i, y_i) we store the pair <distance (r_i) , angle (β_i) > to the reference point (x_{ref}, y_{ref})
- This define the shape regardless its orientation.

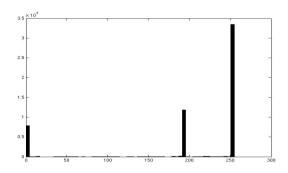
Example:



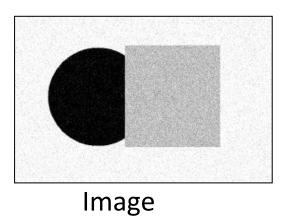
Assumption: different objects present different intensities

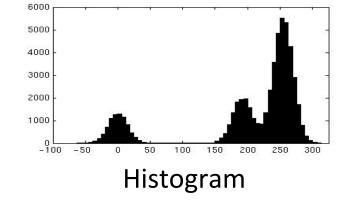
Ideal image (3 objects)





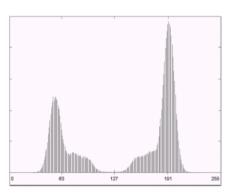
Noisy image (the assumption does not hold)





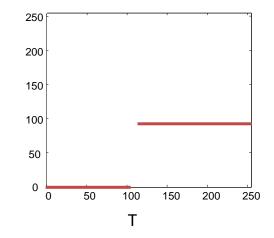
Example:





Objetive: find the threshold T that best separates pixels of similar intensities

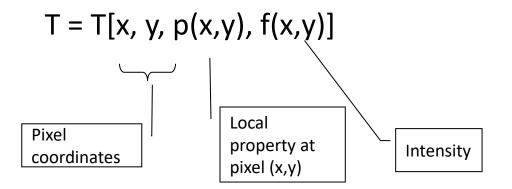
Applying a binarization LUT





Segmented image

The threshold may depend on a number of variables:



$$T = T[f(x,y)]$$

Global threshold (only depends on the intensities)

$$T = T[p(x,y), f(x,y)]$$

Dynamic threshold

$$T = T[x,y, f(x,y)]$$

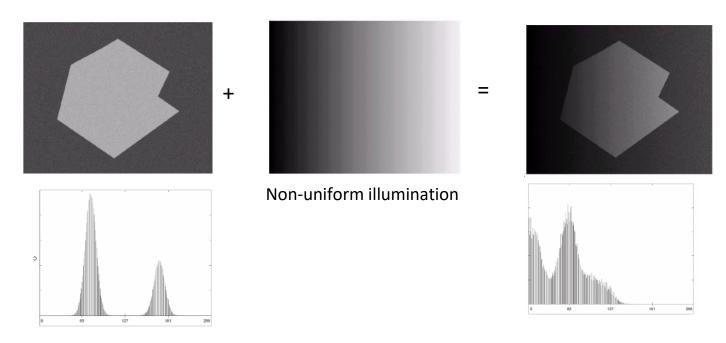
Local threshold

Sometimes, only one threshold is required (i.e. one-color object on a one-color background)



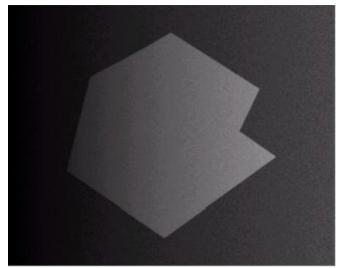
BINARIZATION

Effect of illumination

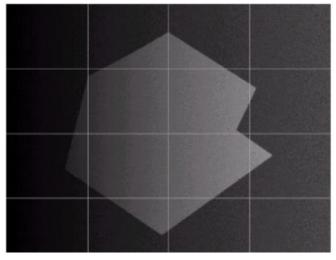


This image can be effectively segmented in two regions by binarization

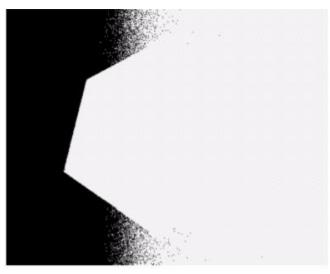
None digitalization will produce good results



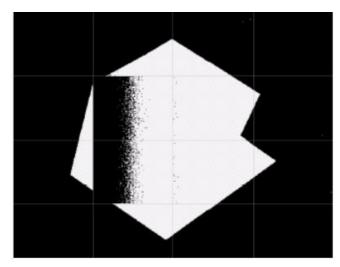
Original Image



Regular partitioning of the image



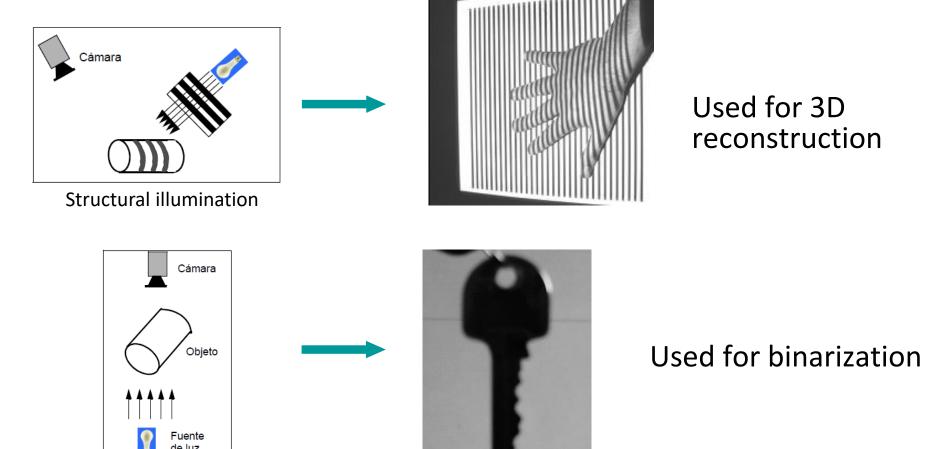
Global threshold (one for the whole image)



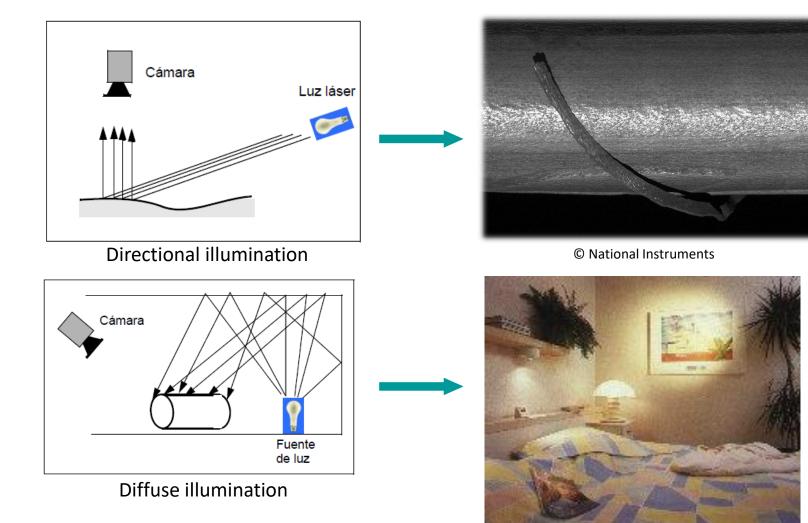
Local threshold (12 different thresholds)

Types of illumination

Back-projection



Types of illumination



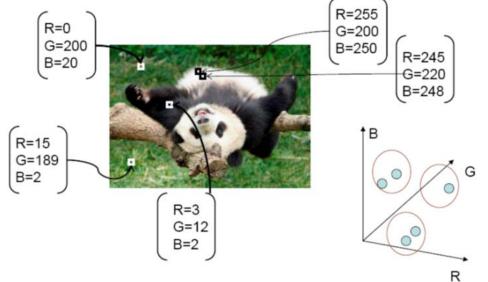
Used for surface inspection

Used for color segmentation

Idea: Group together pixels that are similar according to some properties (*Clustering problem*)

Properties to decide on similarity: intensity, texture, color, pixel location, etc.

Example: property: color (RGB)



Each pixel is a point in the RGB space



Similar pixels are close to each other in this space

Techniques:

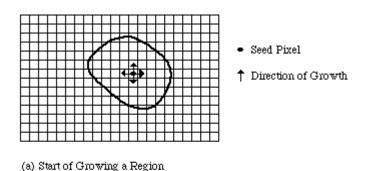
- Region growing
- K-Means
- Expectation-Maximixation
- Mean-Shift

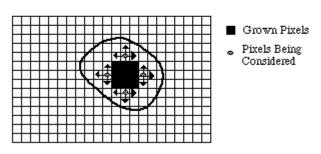
Probabilistic approaches

Source: K. Grauman

Region growing

- Start from a set of seed-pixels which are recursively grown with neighbouring pixels that show similar properties
- If *n* seed-pixels are used, the algorithm ends up with *n* regions, at most (some can be merged)





(b) Growing Process After a Few Iterations

Region growing. Example

Similarity criterion: intensity difference less than 3

Similarity criterion: intensity difference less than 8

0 0 5 6 7
a a b b b
a a b b b
a a b b b
0 1 5 6 5

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0 0 5 6 7
a a a b b
a a a b b
a a a b b
0 1 5 6 5

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Result may depend on the implementation: if the growing process starts from seed 7

0 0 5 6 7
a a b b b
a a b b b
a a b b 5
0 1 5 6 5

a b b b b
a a b b b
a a b b b
a a b b b
a b b b

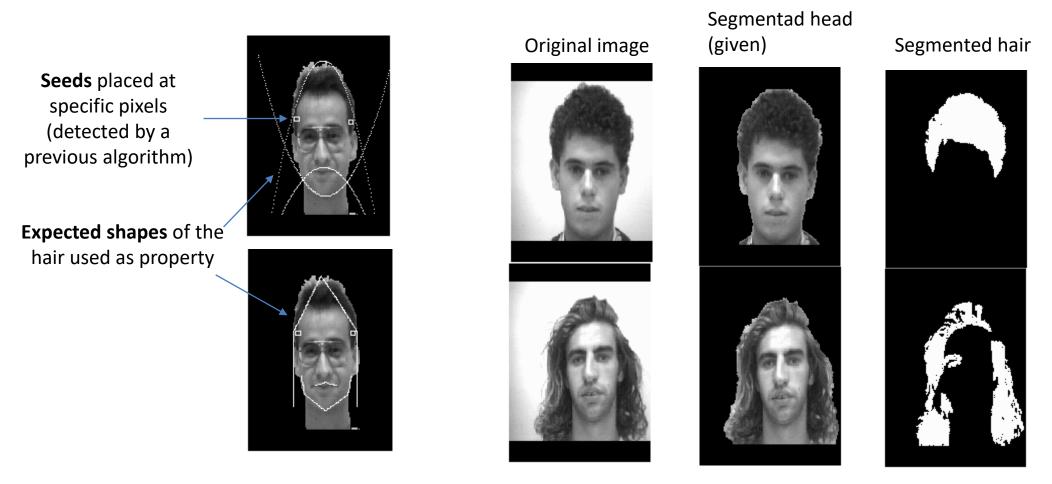
Region growing

Key decisions:

- 1. How many seed pixels
 - Typically, the number of objects we are looking for.
 - Merging of connected regions during the process is possible
- 2. Where to place the seed pixels
 - we must apply any knowledge of the expected regions, if none, at random
- 3. How to select the similarity criterion to add new pixels
 - *Texture*: Images of good resolution are needed.
 - *Color/Intensity*: We can take into account the (dynamic) mean and standard deviation of the current region.
 - **Expected Shape**: We can weight particular directions that more likely fit the expected object shape

4. Region-based techniques Region growing

Example: hair segmentation on a previously segmented head



Property used here: hair pixels must be close to these axes and darker than the face pixels

K-means

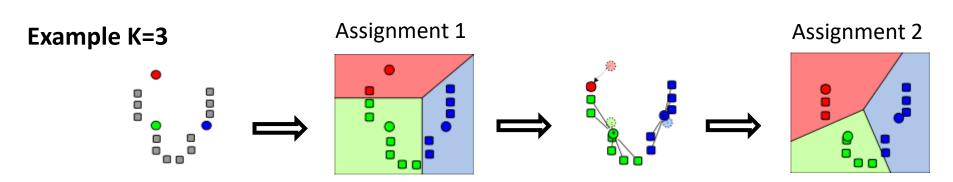
A clustering technique: Given a set of elements, make K clusters out of them

<u>How</u>: minimize the sum of squared Euclidean distances between points x_i and their **nearest cluster centers** m_k

$$\arg\min_{M} D(X, M) = \sum_{\text{cluster } k} \sum_{\text{point } i \text{ in } \atop \text{cluster } k} (x_i - m_k)^2$$

General Algorithm:

- 1. Pick K initial means representing the clusters (not necessary elements of the set)
- 2. Assign each element to the closest mean, so creating new clusters
- 3. Compute the new means of the clusters
- 4. Repeat steps 2 and 3 until convergence

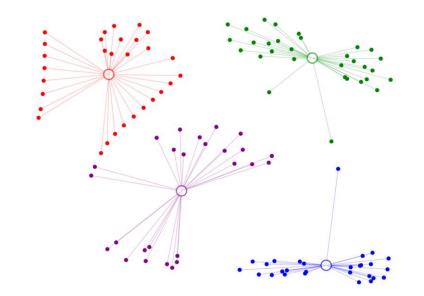


(Wikipedia)

K-means

Similar to region growing, but ...

- All the pixels in the image are classified (assigned to a region) at each step, not just the neighbours
- The algorithm stops when the centers of the regions do not move (in region growing the stop condition is "there are no pixels to add")



Interactive illustration: https://user.ceng.metu.edu.tr/~akifakku s/courses/ceng574/k-means/

K-means for segmentation

- Each pixel of the image is represented by a feature vector (e.g., color, intensity, texture, etc.)
- Each **region** is represented by the **mean of the feature vector** of the pixels in it
- In the feature space we need to define a distance between vectors (e.g. Euclidean distance)

Algorithm

- 1. Select K pixels in the image (manually, at random, with some heuristic, ...) which will represent K regions
- 2. Assign each pixel in the image to the more similar region (the closest one according to the adopted distance)
- 3. Update the feature vector of the new region
- 4. Repeat 2 and 3 until no image pixel changes from one region to another (i.e. region centers do not change) → CONVERGENCE
- 5. Merge connected regions if similar

K-means: Matlab example using intensity as feature vector (1x1)

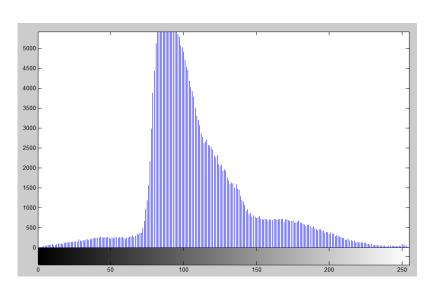
```
im = imread('torre_monica.jpg'); if (size(im,3)==3) im = rgb2gray(im); end
figure, subplot(2,2,1), imshow(im), title('Original Image')
subplot(2,2,3), imhist(im), title('Histogram')
nPixels = prod(size(im)); k=2; % Two classes
data = reshape(im, nPixels,1); %Image as a row vector (one feature)
idx = kmeans(double(data), k); % cluster indices of each pixel (1 or 2)
clust=reshape(idx, size(im)); %Vector-Image back to a matrix
im_clust=uint8(255*clust/max(max(clust)));
subplot(2,2,2), imshow(im_clust), title('Segmented Image')
subplot(2,2,4), imhist(im_clust), title('Histogram')
```

Gray image



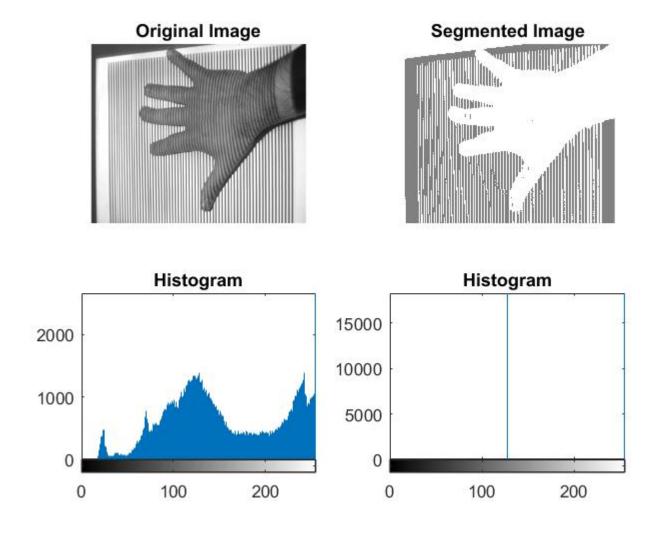
Segmentation from gray



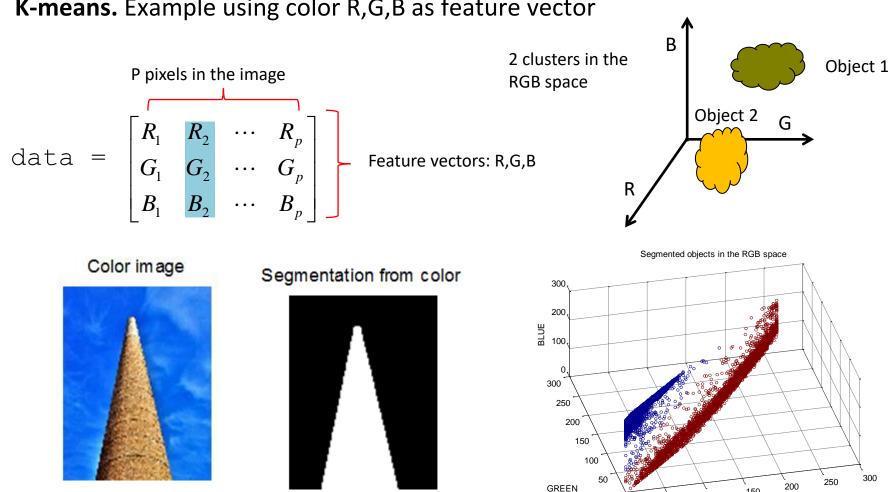


Equivalent to thresholding with the threshold at the middle of the two means (reason why!)

Another example of K-means using intensity as feature vector (dimension 1x1)



K-means. Example using color R,G,B as feature vector



Notice: If the two initial pixels are from the sky the result will not be correct.

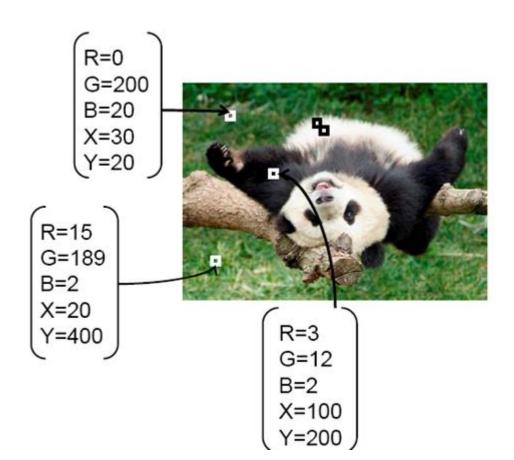
K-means

Usually, color RGB is not enough for a good segmentation



We need some localization feature p.e. (r,g,b,x,y)

Pixel position

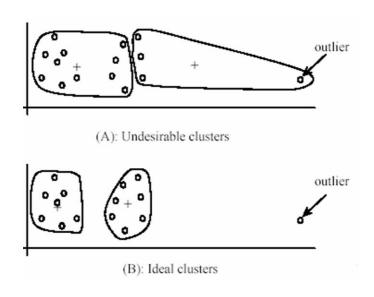


K-means

- Pros
 - Simple
 - Convergence to a local minima (but no guarantee to reach the global minima)

Cons

- High use of memory
- Fixed K
- Sensible to the selection of the initialization
- Sensible to outliers
- Circular clusters in the feature space are assumed (because the use of the Euclidean distance)



Expectation – Maximization (EM)

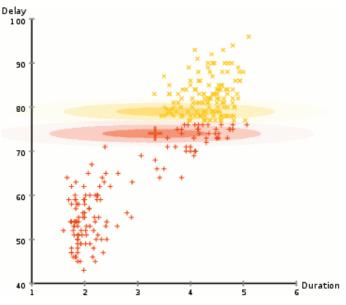
- Is the generalization of the K-means method
- It's a *soft* clustering since it does not give "hard" clusters but the probability that an element (feature vector x of a pixel) belongs to each cluster C_i : $p(x|C_i)$
- Probabilities of the clusters are assumed to be Gaussians: $p(x|C_i) \sim N(\mu_i, \Sigma_i)$

• At each iteration not only the mean is refined (as in K-means), but also the

covariance matrix of the cluster

Example for a two-dimensional feature vector.

Each point is the feature vector of a image pixel



What is a posterior probability?

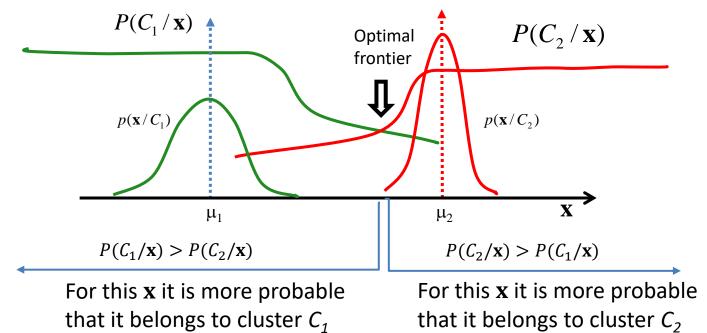
Bayes rule: $P(C_j/\mathbf{x}_i) = \underbrace{\frac{p(\mathbf{x}_i/C_j)P(C_j)}{p(\mathbf{x}_i)}}_{\text{Posterior probability}} = Kp(\mathbf{x}_i/C_j)P(C_j)$ Total probability (does not depend on C_j)

Conditional pdf (o *Likelihood of C_i*) $p(\mathbf{x}_i/C_i)$:

Probability function that a pixel from the cluster C_i has the feature vector \mathbf{x}_i

Prior probability

Example of two clusters:



Pdf: probability density function

Expectation – Maximization

Normal (Gaussian) distribution: Features x_i are continuous random variables that follow the pdf:

$$p(\mathbf{x}_i/C_j) = \frac{1}{(2\pi)^{n/2} |\Sigma_j|^{1/2}} e^{-\frac{1}{2}(\mathbf{x}_i - \mu_j)^T \Sigma_j^{-1} (\mathbf{x}_i - \mu_j)}$$

Feature vector of dimension n: $\mathbf{x} = [x_1 \ x_2 \ \cdots \ x_f \ \cdots \ x_n]^T$

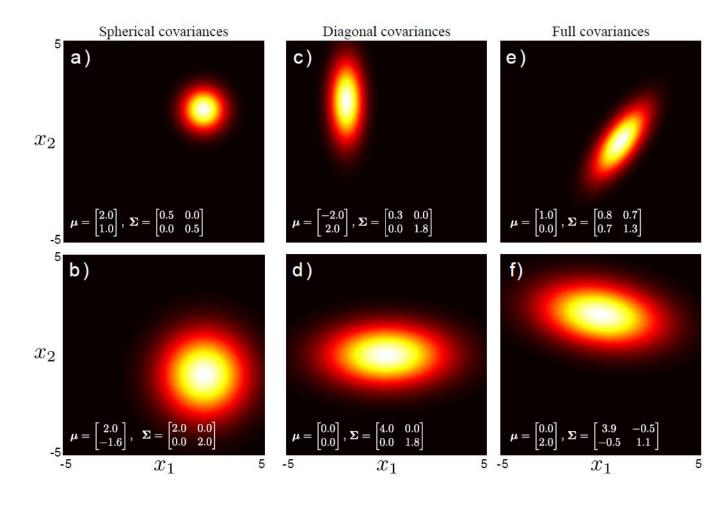
Mean vector: $\mathbf{\mu} = [\mu_1 \quad \mu_2 \quad \cdots \quad \mu_f \quad \cdots \quad \mu_n]^T$

Covariance matrix

$$\Sigma = \mathbb{E}[(\mathbf{x} - \boldsymbol{\mu}) \cdot (\mathbf{x} - \boldsymbol{\mu})^T] = \begin{bmatrix} \sigma_{11} & \cdots & \sigma_{1f} & \cdots & \sigma_{1n} \\ \vdots & & \vdots & & \vdots \\ \sigma_{n1} & \cdots & \sigma_{nf} & \cdots & \sigma_{nn} \end{bmatrix} = \begin{bmatrix} \sigma_{11} & \cdots & 0 & \cdots & 0 \\ \vdots & & \vdots & & \vdots \\ 0 & \cdots & 0 & \cdots & \sigma_{nn} \end{bmatrix}$$
If the features are independents

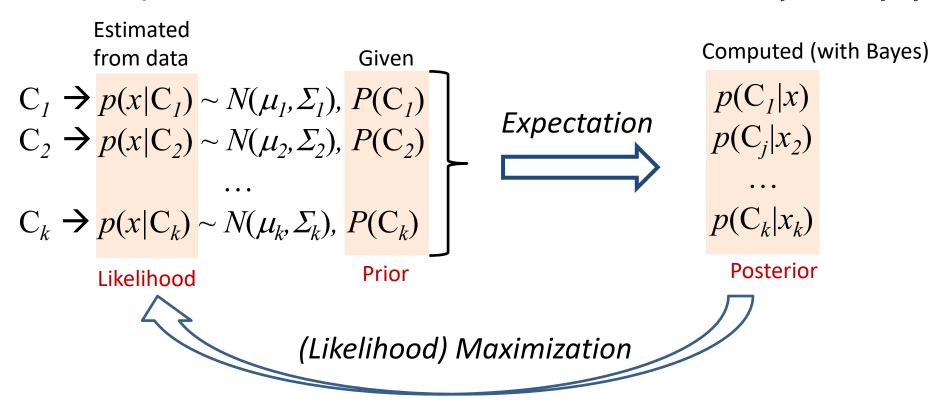
Expectation – Maximization

A Gaussian pdf for different μ and Σ



Expectation – Maximization

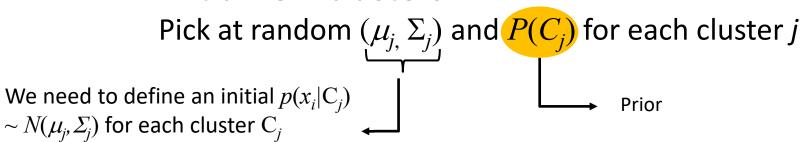
Each cluster C_j has a different and unknown Gaussian pdf $p(x|C_j) \sim N(\mu_j, \Sigma_j)$



If all the $P(C_j)$ are assumed to be identical for all the clusters and $\Sigma_j = \sigma^2 I$ (identical isotropic gaussians), **EM is equivalent to K-means**

EM Algorithm:

Initialize K clusters:



Iteration Steps:

- Using Bayes, compute the cluster C_j to which each data point x_i belongs $p(C_j|x_i)$ Expectation
- Re-estimate the cluster parameters (MLE: Maximum Likelihood estimate)

$$(\mu_j, \Sigma_j), p(C_j)$$
 For each cluster j \longrightarrow Maximization

EM Algorithm:

Iteration steps (until convergence):

Expectation Step: With the current $P(x_i|C_j)$ and $P(C_j)$ compute (via Bayes) the **expected probabilities** $P(C_i|x_i)$ (that x_i belongs to cluster C_i)

$$P(C_{j} | x_{i}) = \frac{p(x_{i} | C_{j}) \cdot p(C_{j})}{p(x_{i})} = \frac{p(x_{i} | C_{j}) \cdot p(C_{j})}{\sum_{i} p(x_{i} | C_{j}) \cdot p(C_{j})}$$

Assign x_i to the cluster C_i with the highest probabilities $P(C_i|x_i)$

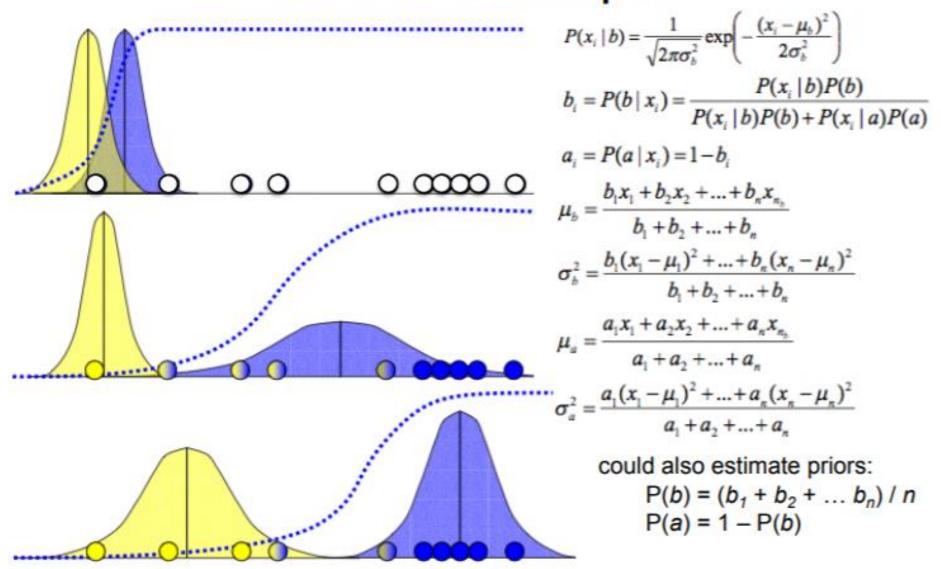
Maximization Step: Estimate $(\mu_j, \Sigma_j), p(C_j)$ that maximize the Likelihood $p(x|C_l)$ (Maximun Likelihood Estimation - MLE)

$$\mu_{j} = \frac{\sum_{i} p(C_{j} \mid x_{i}) \cdot x_{i}}{\sum_{i} p(C_{j} \mid x_{i})} \qquad \Sigma_{j} = \frac{\sum_{i} p(C_{j} \mid x_{i}) \cdot (x_{i} - \mu_{j}) \cdot (x_{i} - \mu_{j})^{T}}{\sum_{i} p(C_{j} \mid x_{i})} \qquad \text{ML estimation of the mean and covariance}$$

 $p(C_j) = \sum_i p(C_j \mid x_i) p(x_i) = \frac{\sum_i p(C_j \mid x_i)}{N}$ If no other information is available, $p(x_i)$ are considered equal-probable

Convergence: when no change ocurrs in a complete iteration

EM: 1-d example



K-means vs. EM

	K-means	EM
Cluster Representation	Mean	Mean, variance
Cluster Initialization	Randomly select K means	Initialize K Gaussian distributions (μ_{j}, Σ_{j}) and $P(C_{j})$
Expectation: Estimate the cluster of each data	Assign each point to the closest mean	Compute $P(C_j x_i)$
Maximization: Re-estimate the cluster parameters	Compute means of current clusters	Compute new $(\mu_{j}, \Sigma_{j}), P(C_{j})$ for each cluster j