# Normalized Cut Method for Image Segmentation

- •J. Shi and J. Malik, *IEEE Trans. Pattern Analysis and Machine Intelligence* **22**(8), 1997
- Divisive (aka splitting, partitioning) method
- Graph-theoretic criterion for measuring goodness of an image partition
- · Hierarchical partitioning
  - dendrogram type representation of all regions

- Criterion for measuring a candidate partitioning:
   Affinity measure between elements within each region is high, and the affinity between elements across regions is low
- Affinity: element x element → R<sup>+</sup> Examples of components of an affinity function: spatial position, intensity, color, texture, motion. Defines the similarity of a pair of data elements.

### **Affinity (Similarity) Measures**

Intensity

$$\operatorname{aff}(\mathbf{x}, \mathbf{y}) = e^{-\|I(\mathbf{x}) - I(\mathbf{y})\|^2 / 2\sigma_I^2}$$

Distance

$$\operatorname{aff}(\mathbf{x}, \mathbf{y}) = e^{-\|\mathbf{x} - \mathbf{y}\|^2 / 2\sigma_d^2}$$

- Color
- Texture
- Motion



#### **Problem Formulation**

- Given an undirected graph G = (V, E), where V is a set of nodes, one for each data element (e.g., pixel), and E is a set of edges with weights representing the affinity between connected nodes
- Find the image partition that maximizes the "association" within each region and minimizes the "disassociation" between regions
- Finding the optimal partition is NP-complete

- Let A, B partition G. Therefore,  $A \cup B = V$ , and  $A \cap B = \emptyset$
- The affinity or similarity between A and B is defined as

$$cut(A,B) = \sum_{i \in A, j \in B} W_{ij}$$
  
= total weight of edges removed

- The **optimal bi-partition** of G is the one that minimizes *cut*
- Cut is biased towards small regions

• Similarly, define the "normalized association:"

$$nassoc(A, B) = \frac{assoc(A, A)}{assoc(A, V)} + \frac{assoc(B, B)}{assoc(B, V)}$$

- Nassoc measures how similar, on average, nodes within the groups are to each other
- New goal: Find the bi-partition that minimizes ncut(A,B) and maximizes nassoc(A,B)
- But, it can be proved that ncut(A,B) = 2 nassoc(A,B), so we can just **minimize** ncut. y = arg min ncut

• So, instead define the **normalized** similarity, called the normalized-cut(A,B), as

$$ncut(A,B) = \frac{cut(A,B)}{assoc(A,V)} + \frac{cut(B,A)}{assoc(B,V)}$$
 where  $assoc(A,V) = \sum_{i \in A, k \in V} W_{ik}$  = total connection weight from nodes in A

to all nodes in G

- Ncut measures the disimilarity between regions ("disassociation" measure)
- Ncut removes the bias based on region size (usually)

• Let y be a P = |V| dimensional vector where

• Let 
$$d(i) = \sum_{i} w_{ij}$$

define the affinity of node *i* with all other nodes

• Let **D** = P x P diagonal matrix:

$$\mathbf{D} = \begin{bmatrix} d_1 & 0 & \dots & 0 \\ 0 & d_2 & \dots & 0 \\ & & \dots & \\ 0 & 0 & \dots & d_{\mathrm{P}} \end{bmatrix}$$
 "degree matrix"

• Let **A** = P x P symmetric matrix:

symmetric matrix: 
$$\mathbf{A} = \begin{bmatrix} w_{11} & w_{12} & \dots & w_{1P} \\ w_{21} & w_{22} & \dots & w_{2P} \\ & & & \dots \\ w_{Pl} & w_{P2} & \dots & w_{PP} \end{bmatrix}$$
 bwn that

 It can be shown that  $y = arg min_x ncut(x)$ 

= 
$$\arg \min_{\mathbf{y}} \frac{\mathbf{y}^{\mathrm{T}}(\mathbf{D} - \mathbf{A})\mathbf{y}}{\mathbf{y}^{\mathrm{T}}\mathbf{D}\mathbf{y}}$$
 subject to  $\mathbf{y}^{\mathrm{T}}\mathbf{D}\mathbf{1} = 0$ 

• Relaxing the constraint on **y** so as to allow it to have real values means that we can approximate the solution by solving an equation of the form:  $(\mathbf{D} - \mathbf{A})\mathbf{y} = \lambda \mathbf{D}\mathbf{y}$ 

- Smallest eigenvector is always 0 because A=V,  $B=\{\}$  means ncut(A,B)=0
- Second smallest eigenvector is the real-valued **y** that minimizes ncut
- Third smallest eigenvector is the real-valued **y** that optimally sub-partitions the first two regions
- Etc.
- Note: Converting from the real-valued **y** to a binaryvalued **y** introduces errors that will propagate to each sub-partition

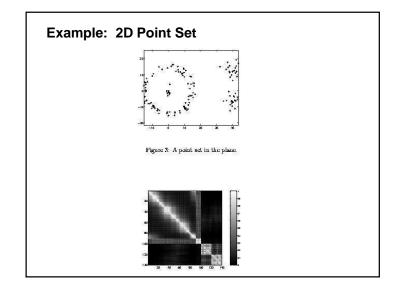
- The solution, y, is an eigenvector of (D A)
- An eigenvector is a characteristic vector of a matrix and specifies a segmentation based on the values of its components; similar points will hopefully have similar eigenvector components.
- Theorem: If **M** is any real, symmetric matrix and **x** is orthogonal to the *j*-1 smallest eigenvectors  $\mathbf{x}_1, \dots, \mathbf{x}_{i+1}$ then  $\mathbf{x}^{\mathsf{T}}\mathbf{M}\mathbf{x} / \mathbf{x}^{\mathsf{T}}\mathbf{x}$  is minimized by the next smallest eigenvector x, and its minimum value is the eigenvalue  $\lambda_i$

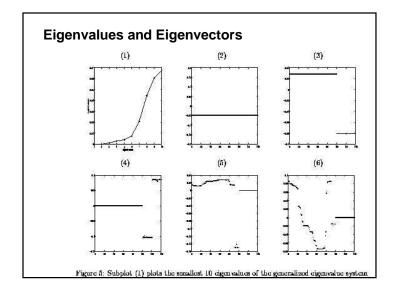
### **NCUT Segmentation Algorithm**

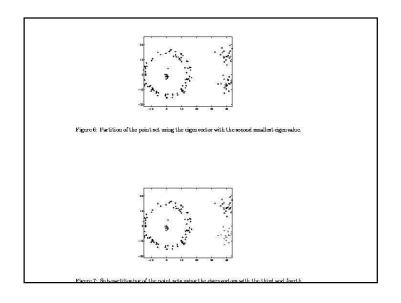
- 1. Set up problem as G = (V, E) and define affinity matrix A and degree matrix D
- 2. Solve  $(\mathbf{D} \mathbf{A})\mathbf{x} = \lambda \mathbf{D}\mathbf{x}$  for the eigenvectors with the smallest eigenvalues
- 3. Let  $\mathbf{x}_2$  = eigenvector with the 2<sup>nd</sup> smallest eigenvalue  $\lambda_2$
- 4. Threshold **x**<sub>2</sub> to obtain the binary-valued vector **x**'<sub>2</sub> such that  $ncut(\mathbf{x'}_2) \ge ncut(\mathbf{x}^{t_2})$  for all possible thresholds t
- 5. For each of the two new regions, if *ncut* < threshold T, then recurse on the region

### **Comments on the Algorithm**

- Recursively bi-partitions the graph instead of using the 3<sup>rd</sup>, 4<sup>th</sup>, etc. eigenvectors for robustness reasons (due to errors caused by the binarization of the real-valued eigenvectors)
- Solving standard eigenvalue problems takes  $O(P^3)$  time
- Can speed up algorithm by exploiting the "locality" of affinity measures, which implies that A is sparse (nonzero values only near the diagonal) and (D − A) is sparse. This leads to a O(P√P) time algorithm







muage segmentation based on brightness and spatial leatures. Figure 8 shows an image that

Example 2: A Grayscale Image



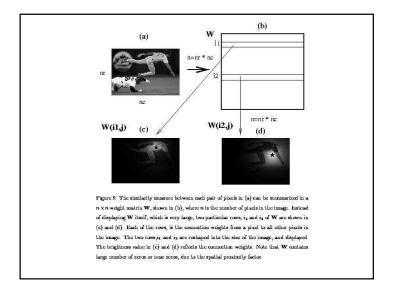
Figure 8: A gray level image of a baseball game.

Just as in the point set grouping case, we have the following steps for image segmentation:

1. Construct a weighted graph, G = (V, E), by taking each pixel as a node, and connecting each pair of pixels by an edge. The weight on that edge should reflect the likelihood of the two pixels belong to one object. Using just the brightness value of the pixels and their

spatial location, we can define the graph edge weight connecting two nodes 
$$i$$
 and  $j$  as:
$$w_{ij} = e^{-\frac{1}{2}\frac{|Y(j-F(j))|^2}{\sigma_d}} * \begin{cases} e^{-\frac{1}{2}\frac{(j-X(j))^2}{\sigma_d}} & \text{if } ||X(i) - X(j)||_2 < r \\ 0 & \text{otherwise} \end{cases}$$
(15)

Figure 9 shows the weight matrix W associated with this weighted graph.



## **Eigenvalues and Eigenvectors**

Putting everything together, each of the matrix-vector computations out O(n) operations with a small constant factor. The number m depends on many factors [11]. In our experiments on image segmentation, we observed that m is typically less than  $O(n^{\frac{1}{2}})$ .

Figure 12 shows the smallest eigenvectors computed for the generalized eigensystem with the weight matrix defined above.

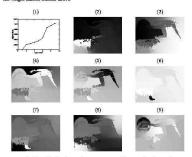


Figure 12: Subplot (i) plots the smallest eigenvectors of the generalized eigenvalue system (ii). Subplot (2) - (9) shows the eigenvectors corresponding the 2nd smallest to the 9th smallest eigenvalues of the system. The eigenvectors are reshaped to be the size of the image.

