Lecture 17: Morphology (ch 7) & Image Matching (ch 13)

ch. 7 and ch. 13 of *Machine Vision* by Wesley E. Snyder & Hairong Qi

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Mathematical Morphology

- The study of shape...
- Using Set Theory
- Most easily understood for binary images.

Binary Morphology: Basic Idea

- 1. Make multiple copies of a shape
- 2. Translate those copies around
- 3. Combine them with either their:
 - Union, U, in the case of dilation, \oplus
 - Intersection, \cap , in the case of erosion, Θ

Dilation makes things bigger Erosion makes things smaller

Binary Morphology: Basic Idea

Q: How do we designate:

- The number of copies to make?
- The translation to apply to each copy?
- A: With a structuring element (s.e.)
 - A (typically) small binary image.
 - We will assume the s.e. always contains the origin.
- For each marked pixel in the s.e.:
 - Make a new copy of the original image
 - Translate that new copy by the coordinates of the current pixel in the s.e.

Dilation Example







 $B = \{ (0,0), (0,-1) \}$

Erosion Example

For erosion, we translate by the *negated* coordinates of the current pixel in the s.e.



Notation

- •A (binary) image: f_A
- The set of marked pixels in f_A : A
 - $\blacksquare \mathsf{A} = \{ (x_1, y_1), (x_2, y_2), \dots \}$
- •A translated image or set: $f_{A_{(dx,dy)}}$ or $A_{(dx,dy)}$
- The number of elements in A: #A
- Complement (inverse) of A: A^c
- Reflection (rotation) of A: Ã

■
$$\tilde{A} = \{ (-x, -y) \mid (x, y) \in A \}$$





Properties

Dilation:

- Commutative, Associative, & Distributive
- Increasing: If $A \subseteq B$ then $A \bigoplus K \subseteq B \bigoplus K$
- Extensive: $A \subseteq A \oplus B$
- Erosion:
 - Anti-extensive (A Θ B \subseteq A), ... (see the text)
- Duality:



Opening

- $f_A \circ f_B = (f_A \Theta f_B) \bigoplus f_B$
- Preserves the geometry of objects that are "big enough"
- Erases smaller objects
- Mental Concept:
 - "Pick up" the s.e. and place it in f_A .
 - Never place the s.e. anywhere it covers any pixels in f_A that are not marked.
 - *f_A* o *f_B* = the set of (marked) pixels in *f_A* which can be covered by the s.e.

Opening Example

•Use a horizontal s.e. to remove 1-pixel thick vertical structures: 1 0





Gray-Scale Morphology

- Morphology operates on sets
- Binary images are just a set of marked pixels
- Gray-scale images contain more information
- How can we apply morphology to this extra intensity information?
- We need to somehow represent intensity as elements of a set

The Umbra

- Gray-scale morphology operates on the umbra of an image.
- Imagine a 2D image as a pixilated surface in 3D
- We can also "pixilate" the height of that surface
- The 2D image is now a 3D surface made of 3D cells







The Distance Transform (DT)

- Records at each pixel the distance from that pixel to the nearest boundary (or to some other feature).
- Used by other algorithms
- The DT is a solution of the Diff. Eq.:
 - $|| \nabla DT(x) || = 1,$
 - DT(x) = 0 on boundary
- Can compute using erosion
 - DT(x) = iteration when x disappears
 - Details in the book

1					1	1	
1					1	1	
1	1	1		1	2	1	
1	2	2	1	2	2	1	
	1	2	2	3	2	1	
	1	2	3	2	2	1	
	1	2	2	1	2	1	
	1	2	1		1	1	
		1			1		
		1					

DT of a region's *interior*

Voronoi Diagram

Divides space

- Related to DT
- Q: To which of a set of regions (or points) is this point the closest?
- Voronoi Diagram's boundaries = points that are equi-distant from multiple regions
- Voronoi Domain of a region = the "cell" of the Voronoi Diagram that contains the region
- Details in the text





Imaging Matching (ch. 13)

- Matching iconic images
- Matching graph-theoretic representations

- Most important:
 - Eigenimages
 - Springs & Templates

Template Matching

- Template ≈ a relatively small reference image for some feature we expect to see in our input image.
- Typical usage: Move the template around the input image, looking for where it "matches" the best (has the highest correlation).
- Rotation & scale can be problematic
 - Often require multiple passes if they can't be ruled out a-priori
- How "big" do we make each template?
 - Do we represent small, simple features
 - Or medium-size, more complex structures?

Eigenimages

- Goal: Identify an image by comparing it to a database of other images
- Problem: Pixel-by-pixel comparisons are two expensive to run across a large database
- Solution: Use PCA



- Big Picture: Fitting a hyper-ellipsoid & then (typically) reducing dimensionality by flattening the shortest axes
- Same as fitting an (N+1)-dimensional multivariate Gaussian, and then taking the level set corresponding to one standard deviation
- Mathematically, PCA reduces the dimensionality of data by mapping it to the first n eigenvectors (*principal components*) of the data's covariance matrix
- The first principal component is the eigenvector with the largest eigenvalue and corresponds to the longest axis of the ellipsoid
- The variance along an eigenvector is exactly the eigenvector's eigenvalue
- This is VERY important and VERY useful. Any questions?

Eigenimages: Procedure

- Run PCA on the training images
 - See the text for efficiency details
- Store in the database:
 - The set of dominant Eigenvectors
 - = the principle components
 - = the Eigenimages
 - For each image, store its coefficients when projected onto the Eigenimages
- Match a new image:
 - Project it onto the basis of the Eigenimages
 - Compare the resulting coefficients to those stored in the database.

Eigenimages Example



The face database and the derived Eigenface examples are all from AT&T Laboratories Cambridge: 20 http://www.cl.cam.ac.uk/research/dtg/attarchive/facedatabase.html & http://en.wikipedia.org/wiki/File:Eigenfaces.png

Matching Simple Features

- Classification based on features
 - Ex: mean intensity, area, aspect ratio
- Idea:
 - Combine a set of shape features into a single feature vector
 - Build a statistical model of this feature vector between and across object classes in a sequence of training shapes
 - Classification of a new shape = the object class from which the new shape's feature vector most likely came.

Graph Matching: Association Graphs

- Match nodes of model to segmented patches in image
- Maximal cliques represent the most likely correspondences
 - Clique = a totally connected subgraph
- Problems: Over/under segmentation, how to develop appropriate rules, often > 1 maximal clique



Graph Matching: Springs & Templates

- Idea: When matching simple templates, we usually expect a certain arrangement between them.
- So, arrange templates using a graph structure.
- The springs are allowed to deform, but only "so" much.



Fischler and Elschlager's "Pictorial Structures" spring & template model for image matching from the early 1970s

Graph Matching: Springs & Templates

- A match is based on minimizing a total cost.
- Problem: Making sure missing a point doesn't improve the score.

 $Cost = \sum_{d \in templates} TemplateCost (d, F(d)) + \sum_{d,e \in ref \times ref} SpringCost (F(d), F(e)) + \sum_{c \in (R_{missing} \times R_{missing})} MissingCost (c)$