

Model Fitting

Introduction to Computer Vision
CSE 152
Lecture 11

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Announcements

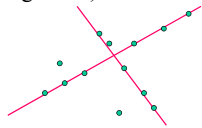
- Homework 3 is due May 10, 11:59 PM
- Reading:
 - Chapter 10: Grouping and Model Fitting

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What to do with edges?

- Segment linked edge chains into curve features (e.g., line segments).
- Group unlinked or unrelated edges into lines (or curves in general).



- Accurately fitting parametric curves (e.g., lines) to grouped edge points.

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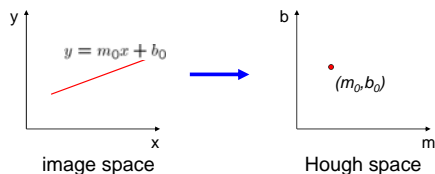
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Hough Transform [Patented 1962]

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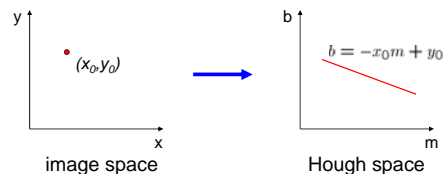
Finding lines in an image



Connection between image (x,y) and Hough (m,b) spaces

- A line in the image corresponds to a point in Hough space

Finding lines in an image



Connection between image (x,y) and Hough (m,b) spaces

- A line in the image corresponds to a point in Hough space
- What does a point (x_0, y_0) in the image space map to?

The equation of any line passing through (x_0, y_0) has form $y_0 = mx_0 + b$

This is a line in Hough space: $b = -x_0m + y_0$

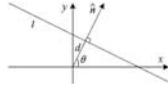
Hough Transform Algorithm

- Typically use a different parameterization

$$d = x \cos \theta + y \sin \theta$$
 - d is the perpendicular distance from the line to the origin
 - θ is the angle this perpendicular makes with the x axis
- Basic Hough transform algorithm
 - Initialize $H[d, \theta] = 0$; H is called accumulator array
 - for each edge point $[x, y]$ in the image
 - for $\theta = 0$ to 180

$$d = x \cos \theta + y \sin \theta$$

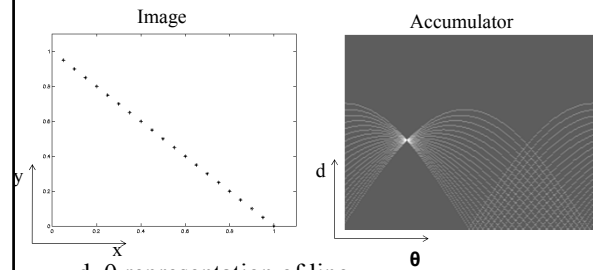
$$H[d, \theta] += 1$$
 - Find the value(s) of (d, θ) where $H[d, \theta]$ is the global maximum
 - The detected line in the image is given by $d = x \cos \theta + y \sin \theta$
- What's the running time (measured in # votes)?



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Hough Transform: 20 colinear points



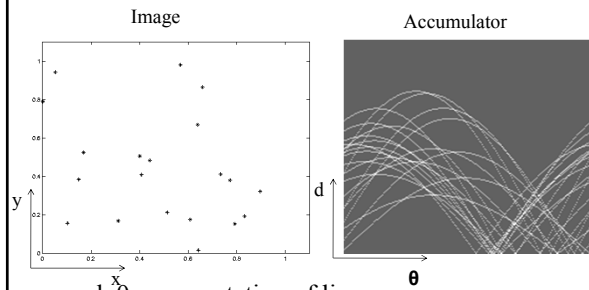
- d, θ representation of line
- Drawn as: $d = |x \cos \theta + y \sin \theta|$
- Maximum accumulator value is 20

Note: accumulator array range: theta: 0-360, d: positive

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Hough Transform: Random points

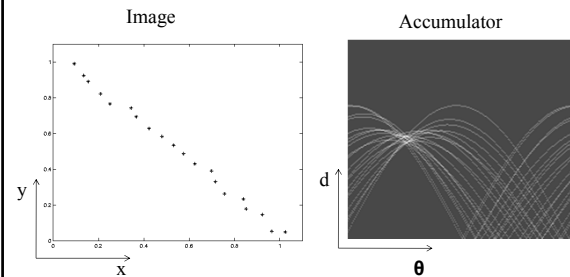


- d, θ representation of line
- Drawn as: $d = |x \cos \theta + y \sin \theta|$
- Maximum accumulator value is 4

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Hough Transform: "Noisy line"



- d, θ representation of line
- Drawn as: $d = |x \cos \theta + y \sin \theta|$
- Maximum accumulator value is 6

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Extension: Oriented Edges

procedure $Hough(\{(x, y, \theta)\})$:

- Clear the accumulator array.
- For each detected edge at location (x, y) and orientation $\theta = \tan^{-1} n_y / n_x$, compute the value of

$$d = x n_x + y n_y$$
 and increment the accumulator corresponding to (θ, d) .
- Find the peaks in the accumulator corresponding to lines.
- Optionally re-fit the lines to the constituent edgels.

Algorithm 4.2 Outline of a Hough transform algorithm based on oriented edge segments.

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Hough Transform for Curves

Generalized Hough Transform

- The Hough transform can be generalized to detect any curve that can be expressed in parametric form:

$$y = f(x, a_1, a_2, \dots, a_p)$$

or

$$g(x, y, a_1, a_2, \dots, a_p) = 0$$

- a_1, a_2, \dots, a_p are the parameters
- The parameter space is p -dimensional
- The accumulating array is *large*

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Example: Finding circles

Equation for circle is

$$(x - x_c)^2 + (y - y_c)^2 = r^2$$

where the parameters are the circle's center (x_c, y_c) and radius r .

Three dimensional generalized Hough space.

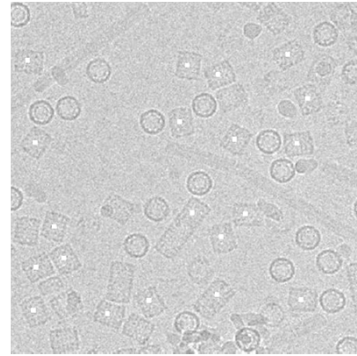
Given an edge point (x, y) ,

1. Loop over all values of (x_c, y_c) ,
2. Compute r
3. Increment $H(x_c, y_c, r)$

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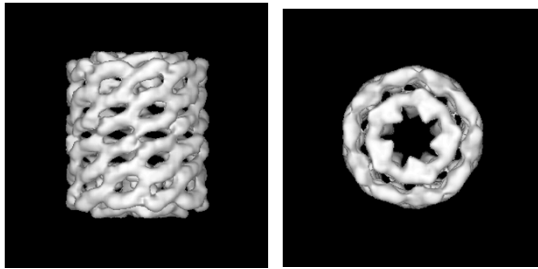
Transmission Electron Microscopy (TEM) Image of Keyhole Limpet Hemocyanin (KLH) with detected particles



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3D Maps of KLH



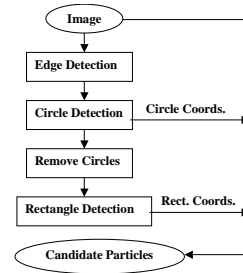
Three-dimensional maps of KLH at a resolution of 23.5 Å reconstructed using particles extracted either manually or automatically as described in the text. (a), (b) The side- and top- view of a 3D map reconstructed from a set of 1042 manually selected particle images.

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Processing in Stage 1 for KLH

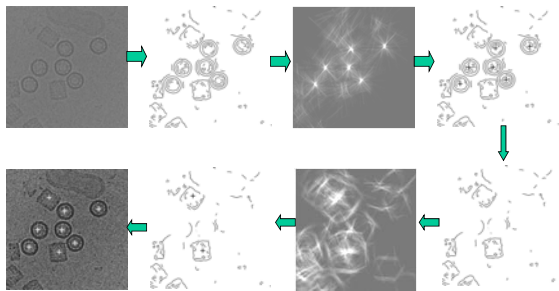
- Canny edge detection.
- A sequence of ordered Hough transforms (HTs) is applied in order from the computationally simplest one to the most complex one.
- Edges covered by the detected shapes are removed immediately from edge images following the application of the last HT.



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Picking KLH Particles in Stage 1

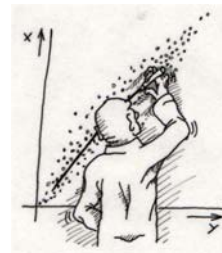


Zhu et al., IEEE Transactions on Medical Imaging, 2003

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Line Fitting



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Line Fitting

Given n points (x_i, y_i) , estimate parameters of line
 $ax_i + by_i - d = 0$
 subject to the constraint that
 $a^2 + b^2 = 1$
 Note: $ax_i + by_i - d$ is distance from (x_i, y_i) to line.

Problem: minimize
 Cost Function:
 Sum of squared distances between each point and the line

$$E(a, b, d) = \sum_{i=1}^n (ax_i + by_i - d)^2$$

with respect to (a, b, d) .

- Minimize E with respect to d :
 $\frac{\partial E}{\partial d} = 0 \Rightarrow d = \frac{1}{n} \sum_{i=1}^n ax_i + by_i = a\bar{x} + b\bar{y}$ Where (\bar{x}, \bar{y}) is the mean of the data points

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Line Fitting

- Substitute d back into E

$$E = \sum_{i=1}^n [a(x_i - \bar{x}) + b(y_i - \bar{y})]^2 = \|U\mathbf{n}\|^2$$
 where $U = \begin{pmatrix} x_1 - \bar{x} & y_1 - \bar{y} \\ \dots & \dots \\ x_n - \bar{x} & y_n - \bar{y} \end{pmatrix}$
 where $\mathbf{n} = (a \ b)^T$.
- Minimize $E = \|U\mathbf{n}\|^2 = \mathbf{n}^T U^T U \mathbf{n} = \mathbf{n}^T S \mathbf{n}$ with respect to a, b subject to the constraint $\mathbf{n}^T \mathbf{n} = 1$. Note that S is given by

$$S = U^T U = \begin{pmatrix} \sum_{i=1}^n x_i^2 - n\bar{x}^2 & \sum_{i=1}^n x_i y_i - n\bar{x}\bar{y} \\ \sum_{i=1}^n x_i y_i - n\bar{x}\bar{y} & \sum_{i=1}^n y_i^2 - n\bar{y}^2 \end{pmatrix}$$

 which is real, symmetric, and positive definite

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Line Fitting

- This is a constrained optimization problem in \mathbf{n} . Solve with Lagrange multiplier

$$L(\mathbf{n}) = \mathbf{n}^T S \mathbf{n} - \lambda(\mathbf{n}^T \mathbf{n} - 1)$$

 Take partial derivative (gradient) w.r.t. \mathbf{n} and set to 0.

$$\nabla L = 2S\mathbf{n} - 2\lambda\mathbf{n} = 0$$

 or

$$S\mathbf{n} = \lambda\mathbf{n}$$

 $\mathbf{n} = (a, b)$ is an Eigenvector of the symmetric matrix S (the one corresponding to the smallest Eigenvalue).
 5. d is computed from Step 1.

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RANSAC

Slides shamelessly taken from Frank Dellaert and Marc Pollefeys and modified

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Simpler Example

- Fitting a straight line

- Inliers
- Outliers

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Discard Outliers

- No point with $d > t$
- RANSAC:
 - RANdom SAMple Consensus
 - Fischler & Bolles 1981
 - Copes with a large proportion of outliers

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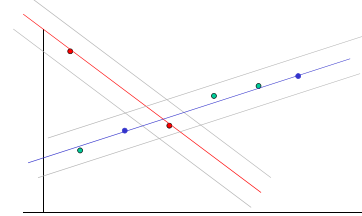
Main Idea

- Select 2 points at random
- Fit a line
- “Support” = number of inliers
- Line with most inliers wins

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Why will this work ?



- Best line has most support
– More support -> better fit

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RANSAC

Objective

Robust fit of model to data set S which contains outliers

Algorithm

- Randomly select a sample of s data points from S and instantiate the model from this subset.
- Determine the set of data points S_i which are within a distance threshold t of the model. The set S_i is the **consensus set** of samples and defines the inliers of S .
- If the size of S_i is greater than some threshold T , re-estimate the model using all the points in S_i and terminate
- If the size of S_i is less than T , select a new subset and repeat the above.
- After N trials the largest consensus set S_i is selected, and the model is re-estimated using all the points in the subset S_i .

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Number of trials

Choose N (number of trials) so that, with probability p , at least one random sample is free from outliers. e.g., $p=0.99$

$$(1 - (1 - e)^s)^N = 1 - p$$

$$N = \log(1 - p) / \log(1 - (1 - e)^s)$$

e : proportion of outliers

s : Number of points needed for the model

s	proportion of outliers e						
	5%	10%	20%	25%	30%	40%	50%
2	2	3	5	6	7	11	17
3	3	4	7	9	11	19	35
4	3	5	9	13	17	34	72
5	4	6	12	17	26	57	146
6	4	7	16	24	37	97	293
7	4	8	20	33	54	163	588
8	5	9	26	44	78	272	1177

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Number of inliers threshold

- Typically, terminate when inlier ratio reaches expected ratio of inliers

$$T = (1 - e)N$$

- N (number of trials)
- e proportion of outliers

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Distance threshold

Choose threshold t so probability for inlier is a (e.g., 0.95)

- Often empirically
- Zero-mean Gaussian noise σ then d_{\perp}^2 follows χ_m^2 distribution with m =codimension of model
(codimension=dimension of space – dimension of subspace)

Codimension	Model	t^2
1	E, F, 2D line	$3.84\sigma^2$
2	P	$5.99\sigma^2$

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Using RANSAC to estimate the Fundamental Matrix

- What is the model?
- What is the sample size and where do the samples come from?
- What distance do we use to compute the consensus set?
- How often do outliers occur?

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Other models

- 2D motion models
- Typically: points in two images
- Candidates:
 - Translation
 - Euclidean
 - Similarity
 - Affine
 - Projective

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Feature Detection and Matching

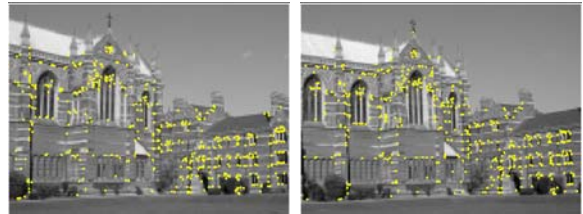


Input Images

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Feature Detection and Matching

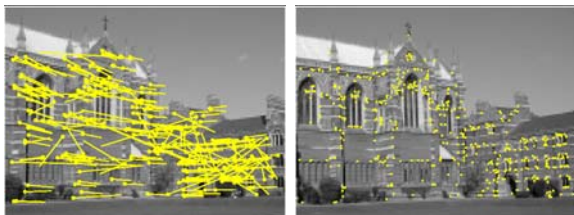


Detected Corners

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Feature Detection and Matching

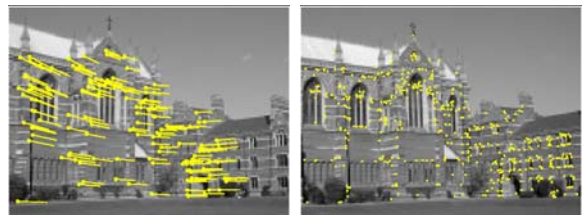


Simple Matching

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Feature Detection and Matching



Simple Matching
Including Outlier Rejection

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Mosaicing: Homography Estimate with RANSAC



www.cs.cmu.edu/~dellaert/mosaicking

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Next Lecture

- Motion
- Reading:
 - Section 10.6.1: Optical Flow and Motion
 - Section 10.6.2: Flow Models
 - Introductory Techniques for 3-D Computer Vision, Trucco and Verri
 - Chapter 8: Motion

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