



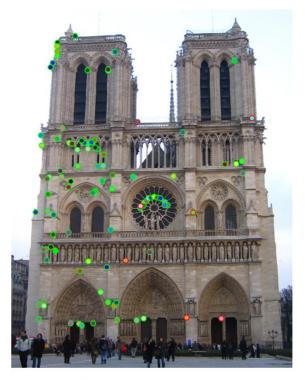
Fitting and Alignment

Szeliski 6.1

Computer Vision CS 143, Brown

James Hays

Project 2 questions?





The top 100 most confident local feature matches from a baseline implementation of project 2. In this case, 93 were correct (highlighted in green) and 7 were incorrect (highlighted in red).

Project 2: Local Feature Matching

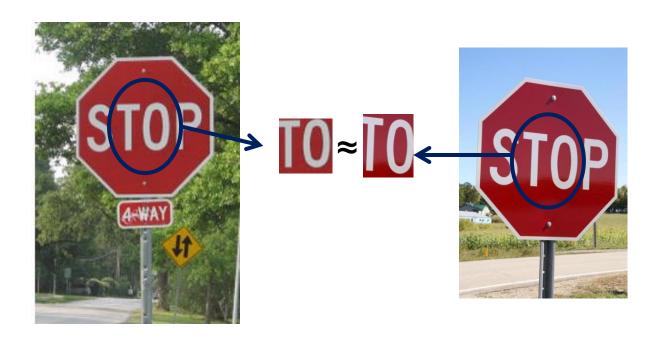
CS 143: Introduction to Computer Vision

Brief

- · Due: 11:59pm on Monday, October 7th, 2013
- Stencil code: /course/cs143/asgn/proj2/code/
- Data: /course/cs143/asqn/proj2/data/ includes 93 images from 9 different outdoor scenes.
- Html writeup template: /course/cs143/asgn/proj2/html/
- · Partial project materials are also available in proj2.zip (1.7 MB). Includes only the two test images shown above.
- Handin: cs143 handin proj2
- · Required files: README, code/, html/, html/index.html

This section: correspondence and alignment

 Correspondence: matching points, patches, edges, or regions across images



Review

Fitting: find the parameters of a model that best fit the data

Alignment: find the parameters of the transformation that best align matched points

Review: Fitting and Alignment

- Design challenges
 - Design a suitable goodness of fit measure
 - Similarity should reflect application goals
 - Encode robustness to outliers and noise
 - Design an optimization method
 - Avoid local optima
 - Find best parameters quickly

Fitting and Alignment: Methods

- Global optimization / Search for parameters
 - Least squares fit
 - Robust least squares
 - Iterative closest point (ICP)

- Hypothesize and test
 - Hough transform
 - RANSAC

Review: Hough Transform

1. Create a grid of parameter values

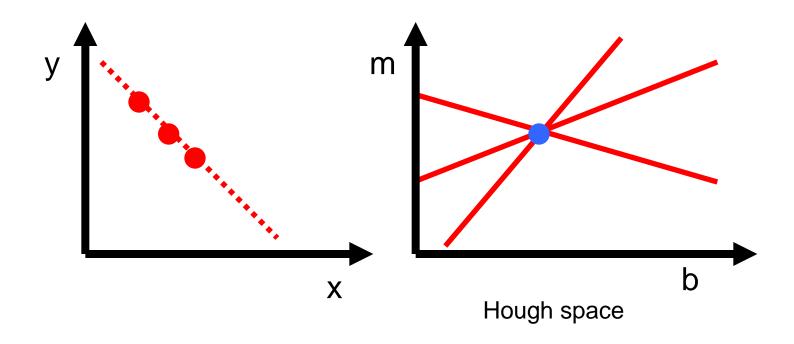
2. Each point votes for a set of parameters, incrementing those values in grid

3. Find maximum or local maxima in grid

Review: Hough transform

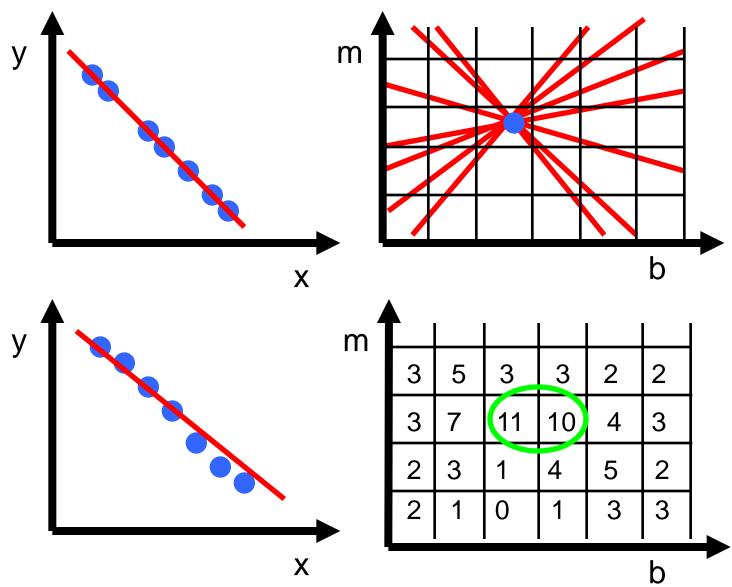
P.V.C. Hough, *Machine Analysis of Bubble Chamber Pictures,* Proc. Int. Conf. High Energy Accelerators and Instrumentation, 1959

Given a set of points, find the curve or line that explains the data points best



$$y = m x + b$$

Review: Hough transform



Hough Transform

- How would we find circles?
 - Of fixed radius
 - Of unknown radius

Hough transform conclusions

Good

- Robust to outliers: each point votes separately
- Fairly efficient (much faster than trying all sets of parameters)
- Provides multiple good fits

Bad

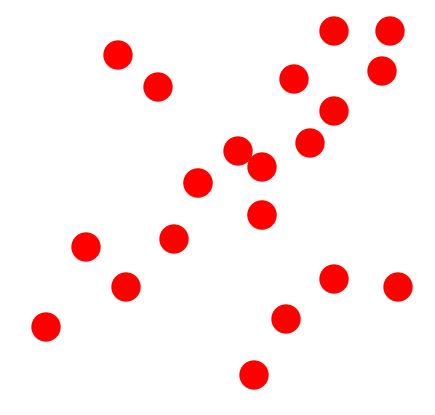
- Some sensitivity to noise
- Bin size trades off between noise tolerance, precision, and speed/memory
 - Can be hard to find sweet spot
- Not suitable for more than a few parameters
 - grid size grows exponentially

Common applications

- Line fitting (also circles, ellipses, etc.)
- Object instance recognition (parameters are affine transform)
- Object category recognition (parameters are position/scale)

(RANdom SAmple Consensus):

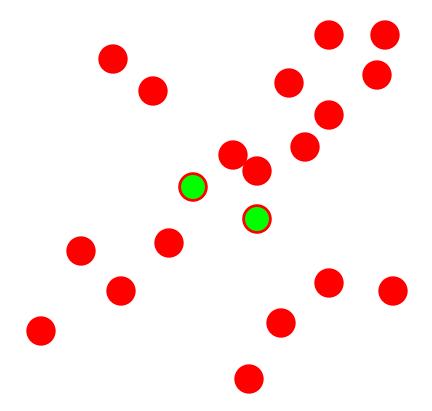
Fischler & Bolles in '81.



Algorithm:

- 1. **Sample** (randomly) the number of points required to fit the model
- 2. **Solve** for model parameters using samples
- 3. **Score** by the fraction of inliers within a preset threshold of the model

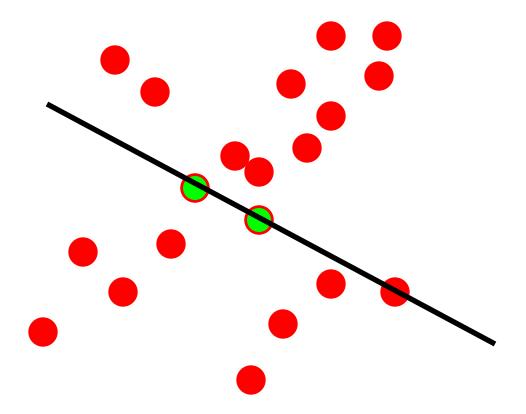
Line fitting example



Algorithm:

- 1. **Sample** (randomly) the number of points required to fit the model (#=2)
- 2. **Solve** for model parameters using samples
- 3. **Score** by the fraction of inliers within a preset threshold of the model

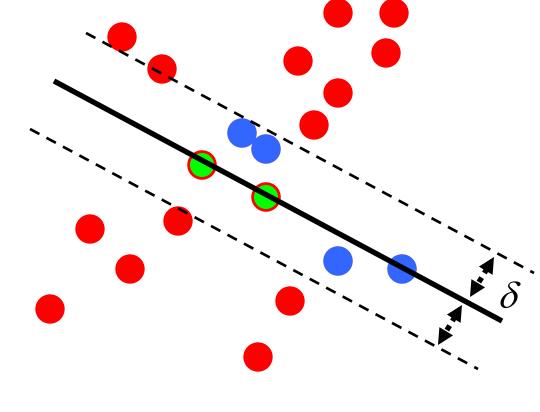
Line fitting example



Algorithm:

- 1. Sample (randomly) the number of points required to fit the model (#=2)
- 2. **Solve** for model parameters using samples
- 3. **Score** by the fraction of inliers within a preset threshold of the model

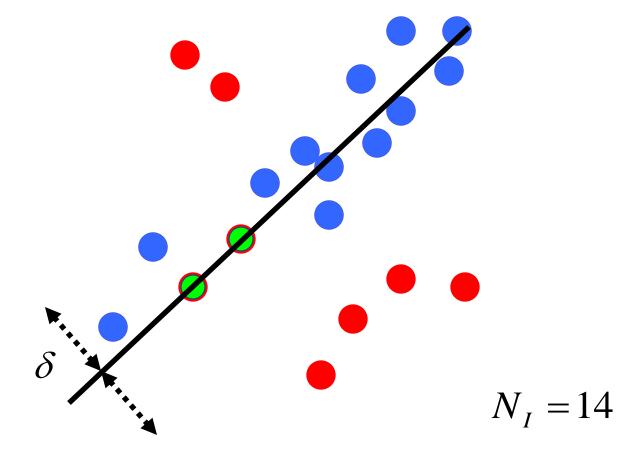
Line fitting example



$$N_I = 6$$

Algorithm:

- Sample (randomly) the number of points required to fit the model (#=2)
- 2. Solve for model parameters using samples
- 3. **Score** by the fraction of inliers within a preset threshold of the model



Algorithm:

- 1. **Sample** (randomly) the number of points required to fit the model (#=2)
- 2. **Solve** for model parameters using samples
- 3. **Score** by the fraction of inliers within a preset threshold of the model

How to choose parameters?

- Number of samples N
 - Choose N so that, with probability p, at least one random sample is free from outliers (e.g. p=0.99) (outlier ratio: e)
- Number of sampled points s
 - Minimum number needed to fit the model
- Distance threshold δ
 - Choose δ so that a good point with noise is likely (e.g., prob=0.95) within threshold
 - Zero-mean Gaussian noise with std. dev. σ : $t^2=3.84\sigma^2$

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

	proportion of outliers e							
S	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	

RANSAC conclusions

Good

- Robust to outliers
- Applicable for larger number of objective function parameters than Hough transform
- Optimization parameters are easier to choose than Hough transform

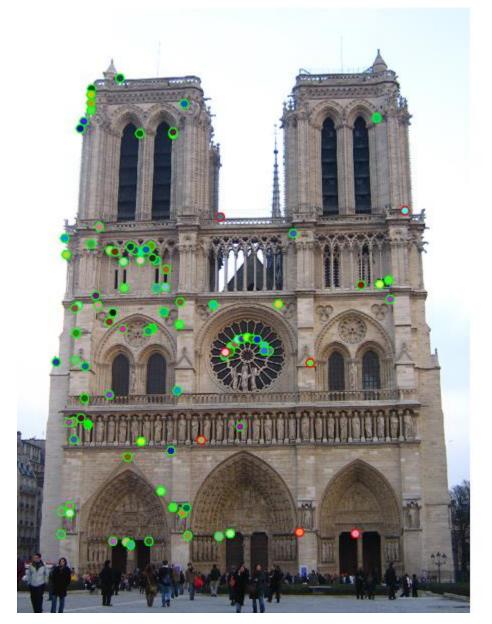
Bad

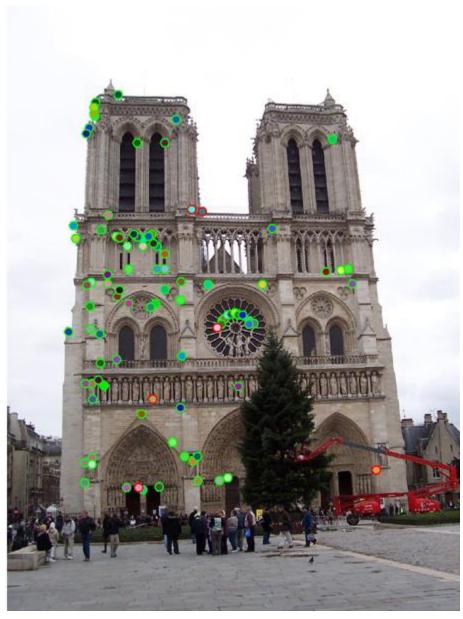
- Computational time grows quickly with fraction of outliers and number of parameters
- Not good for getting multiple fits

Common applications

- Computing a homography (e.g., image stitching)
- Estimating fundamental matrix (relating two views)

How do we fit the best alignment?





Alignment

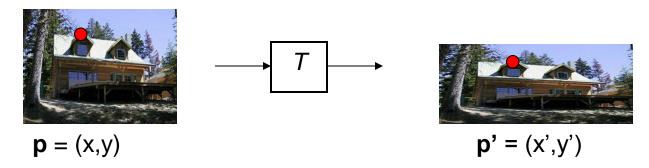
 Alignment: find parameters of model that maps one set of points to another

 Typically want to solve for a global transformation that accounts for *most* true correspondences

Difficulties

- Noise (typically 1-3 pixels)
- Outliers (often 50%)
- Many-to-one matches or multiple objects

Parametric (global) warping



Transformation T is a coordinate-changing machine:

$$p' = T(p)$$

What does it mean that T is global?

- Is the same for any point p
- can be described by just a few numbers (parameters)

For linear transformations, we can represent T as a matrix

$$p' = Tp$$

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \mathbf{T} \begin{bmatrix} x \\ y \end{bmatrix}$$

Common transformations



original

Transformed



translation



rotation



aspect



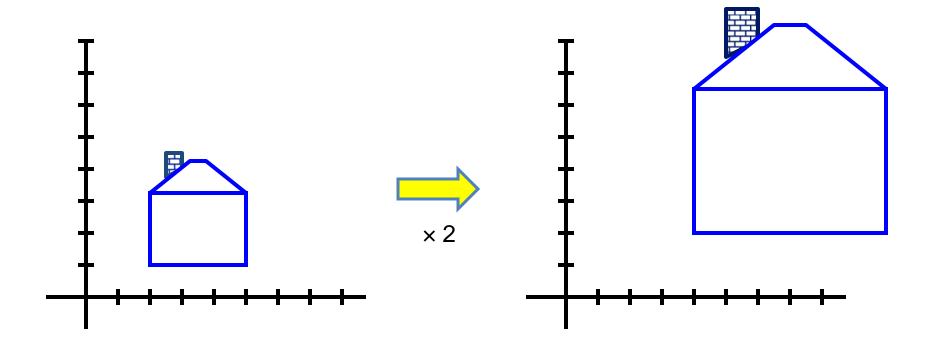
affine



perspective

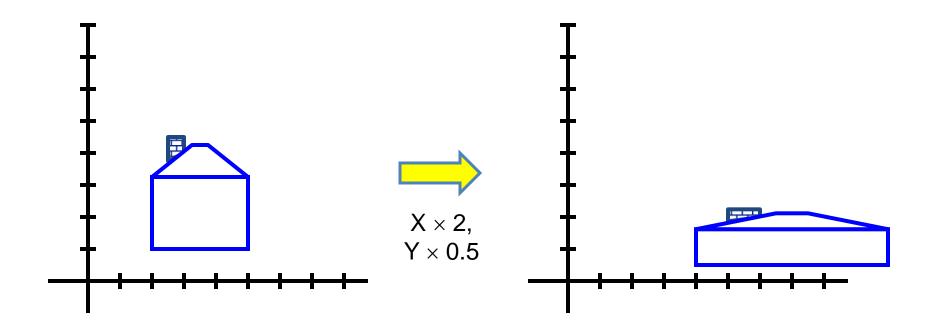
Scaling

- Scaling a coordinate means multiplying each of its components by a scalar
- *Uniform scaling* means this scalar is the same for all components:



Scaling

• *Non-uniform scaling*: different scalars per component:



Scaling

Scaling operation:

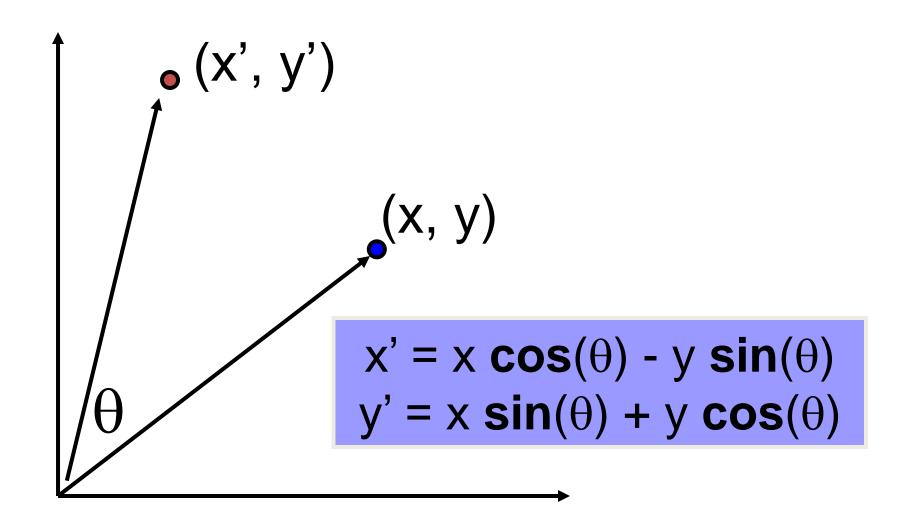
$$x' = ax$$

$$y' = by$$

• Or, in matrix form:

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} a & 0 \\ 0 & b \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix}$$
scaling matrix S

2-D Rotation



2-D Rotation

This is easy to capture in matrix form:

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} \cos(\theta) & -\sin(\theta) \\ \sin(\theta) & \cos(\theta) \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix}$$

Even though $sin(\theta)$ and $cos(\theta)$ are nonlinear functions of θ ,

- -x' is a linear combination of x and y
- -y' is a linear combination of x and y

What is the inverse transformation?

- Rotation by $-\theta$
- For rotation matrices $\mathbf{R}^{-1} = \mathbf{R}^T$

Basic 2D transformations

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} s_x & 0 \\ 0 & s_y \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix}$$
Scale

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} \cos\Theta & -\sin\Theta \\ \sin\Theta & \cos\Theta \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix}$$

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} 1 & \alpha_x \\ \alpha_y & 1 \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix}$$
Shear

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} 1 & 0 & t_x \\ 0 & 1 & t_y \end{bmatrix} \begin{bmatrix} x \\ y \\ 1 \end{bmatrix}$$
Translate

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \end{bmatrix} \begin{bmatrix} x \\ y \\ 1 \end{bmatrix}$$
Affine

 $\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \end{bmatrix} \begin{vmatrix} x \\ y \\ 1 \end{vmatrix}$ Affine is any combination of translation, scale, rotation, shear

Affine Transformations

Affine transformations are combinations of

- Linear transformations, and
- Translations

Properties of affine transformations:

- Lines map to lines
- Parallel lines remain parallel
- Ratios are preserved
- Closed under composition

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \end{bmatrix} \begin{bmatrix} x \\ y \\ 1 \end{bmatrix}$$

or

$$\begin{bmatrix} x' \\ y' \\ 1 \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} x \\ y \\ 1 \end{bmatrix}$$

Projective Transformations

Projective transformations are combos of

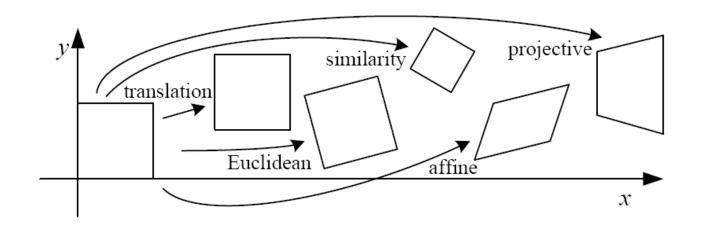
- Affine transformations, and
- Projective warps

$$\begin{bmatrix} x' \\ y' \\ w' \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \\ g & h & i \end{bmatrix} \begin{bmatrix} x \\ y \\ w \end{bmatrix}$$

Properties of projective transformations:

- Lines map to lines
- Parallel lines do not necessarily remain parallel
- Ratios are not preserved
- Closed under composition
- Models change of basis
- Projective matrix is defined up to a scale (8 DOF)

2D image transformations (reference table)

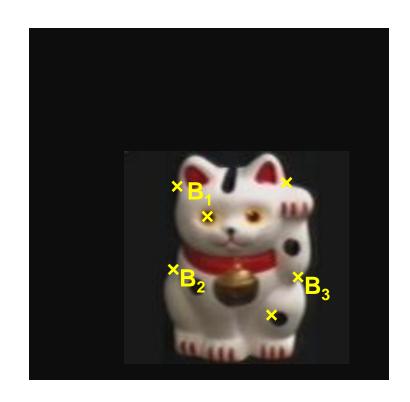


Name	Matrix	# D.O.F.	Preserves:	Icon
translation	$egin{bmatrix} ig[egin{array}{c c} I & t \end{bmatrix}_{2 imes 3} \end{array}$	2	orientation $+ \cdots$	
rigid (Euclidean)	$\left[egin{array}{c c} R & t\end{array} ight]_{2 imes 3}$	3	lengths + · · ·	
similarity	$\left[\begin{array}{c c} sR \mid t\end{array}\right]_{2 imes 3}$	4	angles $+\cdots$	
affine	$\left[egin{array}{c} oldsymbol{A} \end{array} ight]_{2 imes 3}$	6	parallelism + · · ·	
projective	$\left[egin{array}{c} ilde{m{H}} \end{array} ight]_{3 imes 3}$	8	straight lines	

Szeliski 2.1

Example: solving for translation

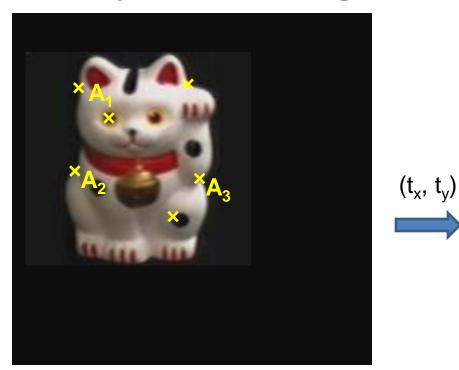


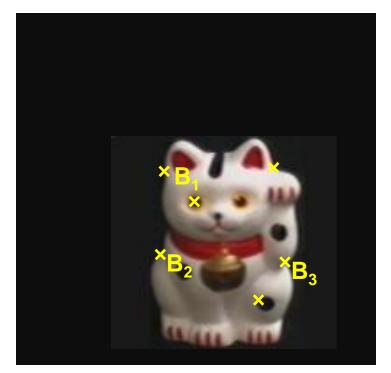


Given matched points in {A} and {B}, estimate the translation of the object

$$\begin{bmatrix} x_i^B \\ y_i^B \end{bmatrix} = \begin{bmatrix} x_i^A \\ y_i^A \end{bmatrix} + \begin{bmatrix} t_x \\ t_y \end{bmatrix}$$

Example: solving for translation



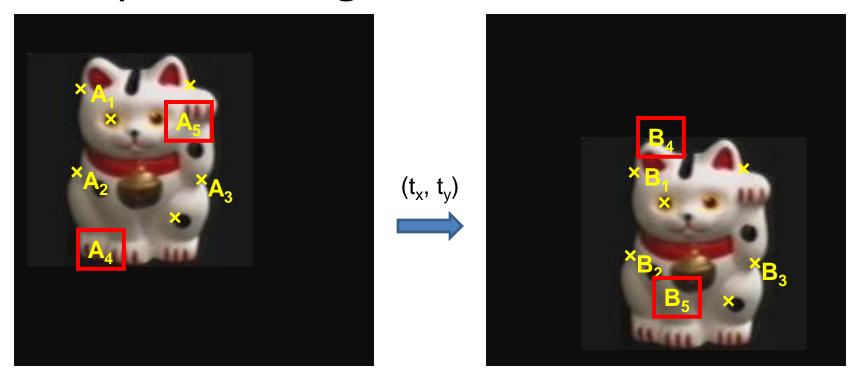


Least squares solution

- 1. Write down objective function
- 2. Derived solution
 - a) Compute derivative
 - b) Compute solution
- 3. Computational solution
 - a) Write in form Ax=b
 - b) Solve using pseudo-inverse or eigenvalue decomposition

$$\begin{bmatrix} x_i^B \\ y_i^B \end{bmatrix} = \begin{bmatrix} x_i^A \\ y_i^A \end{bmatrix} + \begin{bmatrix} t_x \\ t_y \end{bmatrix}$$

$$\begin{bmatrix} 1 & 0 \\ 0 & 1 \\ \vdots & \vdots \\ 1 & 0 \\ 0 & 1 \end{bmatrix} \begin{bmatrix} t_x \\ t_y \end{bmatrix} = \begin{bmatrix} x_1^B - x_1^A \\ y_1^B - y_1^A \\ \vdots \\ x_n^B - x_n^A \\ y_n^B - y_n^A \end{bmatrix}$$

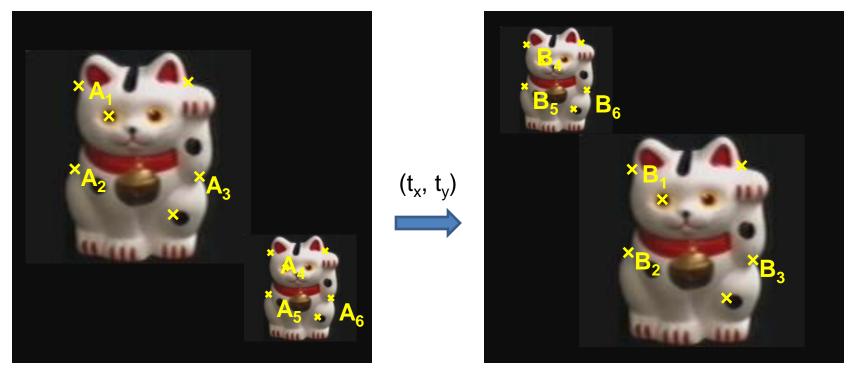


Problem: outliers

RANSAC solution

- 1. Sample a set of matching points (1 pair)
- 2. Solve for transformation parameters
- 3. Score parameters with number of inliers
- 4. Repeat steps 1-3 N times

$$\begin{bmatrix} x_i^B \\ y_i^B \end{bmatrix} = \begin{bmatrix} x_i^A \\ y_i^A \end{bmatrix} + \begin{bmatrix} t_x \\ t_y \end{bmatrix}$$

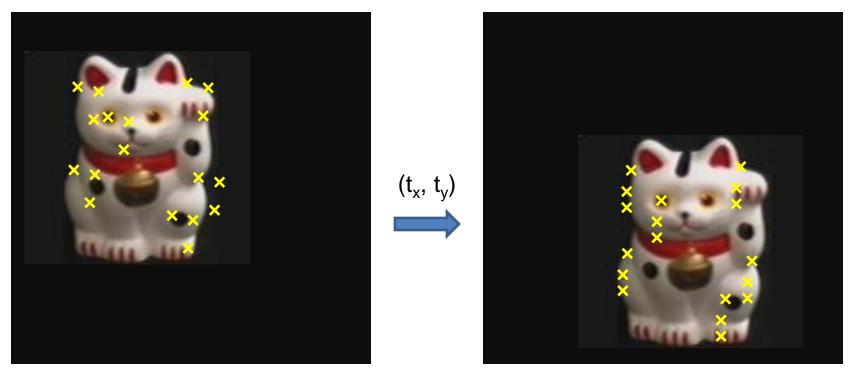


Problem: outliers, multiple objects, and/or many-to-one matches

Hough transform solution

- 1. Initialize a grid of parameter values
- 2. Each matched pair casts a vote for consistent values
- 3. Find the parameters with the most votes
- 4. Solve using least squares with inliers

$$\begin{bmatrix} x_i^B \\ y_i^B \end{bmatrix} = \begin{bmatrix} x_i^A \\ y_i^A \end{bmatrix} + \begin{bmatrix} t_x \\ t_y \end{bmatrix}$$



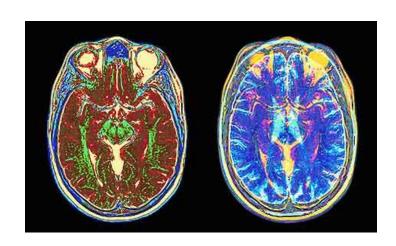
Problem: no initial guesses for correspondence

$$\begin{bmatrix} x_i^B \\ y_i^B \end{bmatrix} = \begin{bmatrix} x_i^A \\ y_i^A \end{bmatrix} + \begin{bmatrix} t_x \\ t_y \end{bmatrix}$$

What if you want to align but have no prior matched pairs?

Hough transform and RANSAC not applicable

Important applications



Medical imaging: match brain scans or contours



Robotics: match point clouds

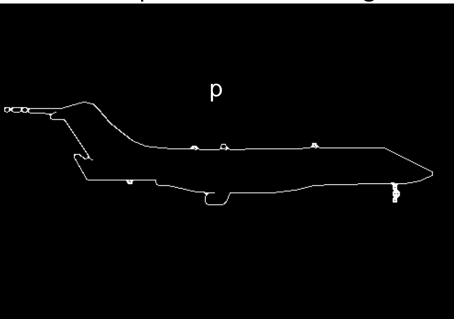
Iterative Closest Points (ICP) Algorithm

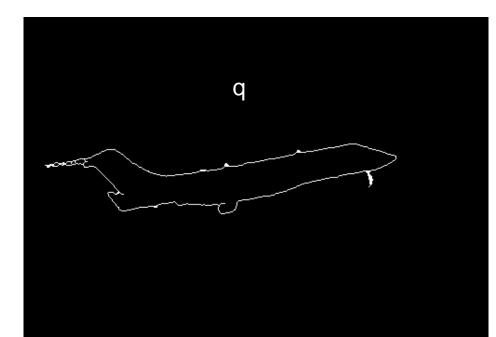
Goal: estimate transform between two dense sets of points

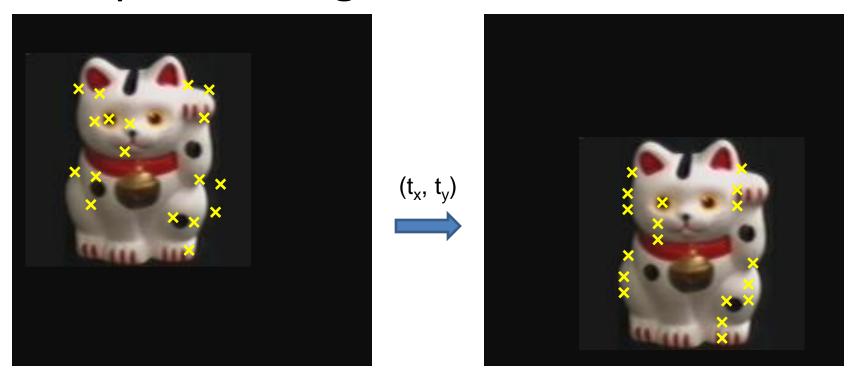
- Initialize transformation (e.g., compute difference in means and scale)
- **2. Assign** each point in {Set 1} to its nearest neighbor in {Set 2}
- 3. Estimate transformation parameters
 - e.g., least squares or robust least squares
- **4. Transform** the points in {Set 1} using estimated parameters
- **5. Repeat** steps 2-4 until change is very small

Example: aligning boundaries

- 1. Extract edge pixels $p_1 \dots p_n$ and $q_1 \dots q_m$
- 2. Compute initial transformation (e.g., compute translation and scaling by center of mass, variance within each image)
- 3. Get nearest neighbors: for each point p_i find corresponding match(i) = $\underset{j}{\operatorname{argmin}} dist(pi, qj)$
- 4. Compute transformation **T** based on matches
- 5. Warp points **p** according to **T**
- 6. Repeat 3-5 until convergence







Problem: no initial guesses for correspondence

ICP solution

- 1. Find nearest neighbors for each point
- 2. Compute transform using matches
- 3. Move points using transform
- 4. Repeat steps 1-3 until convergence

$$\begin{bmatrix} x_i^B \\ y_i^B \end{bmatrix} = \begin{bmatrix} x_i^A \\ y_i^A \end{bmatrix} + \begin{bmatrix} t_x \\ t_y \end{bmatrix}$$

Algorithm Summary

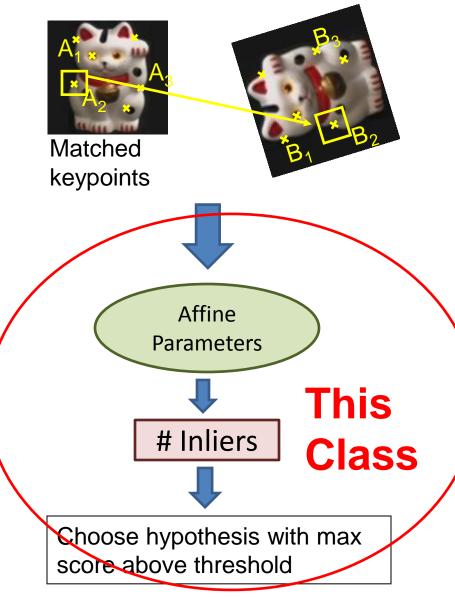
- Least Squares Fit
 - closed form solution
 - robust to noise
 - not robust to outliers
- Robust Least Squares
 - improves robustness to noise
 - requires iterative optimization
- Hough transform
 - robust to noise and outliers
 - can fit multiple models
 - only works for a few parameters (1-4 typically)
- RANSAC
 - robust to noise and outliers
 - works with a moderate number of parameters (e.g, 1-8)
- Iterative Closest Point (ICP)
 - For local alignment only: does not require initial correspondences

Object Instance Recognition

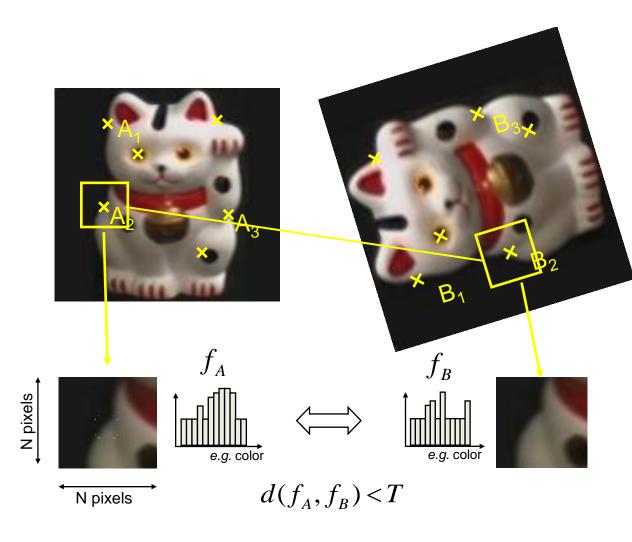
1. Match keypoints to object model

2. Solve for affine transformation parameters

3. Score by inliers and choose solutions with score above threshold

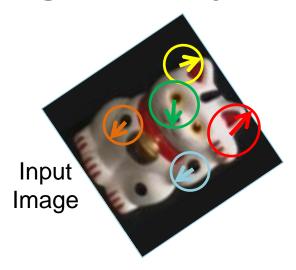


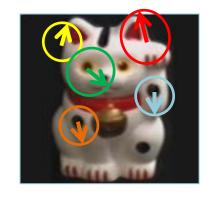
Overview of Keypoint Matching



- 1. Find a set of distinctive key-points
- 2. Define a region around each keypoint
- 3. Extract and normalize the region content
- 4. Compute a local descriptor from the normalized region
- 5. Match local descriptors

Finding the objects (overview)



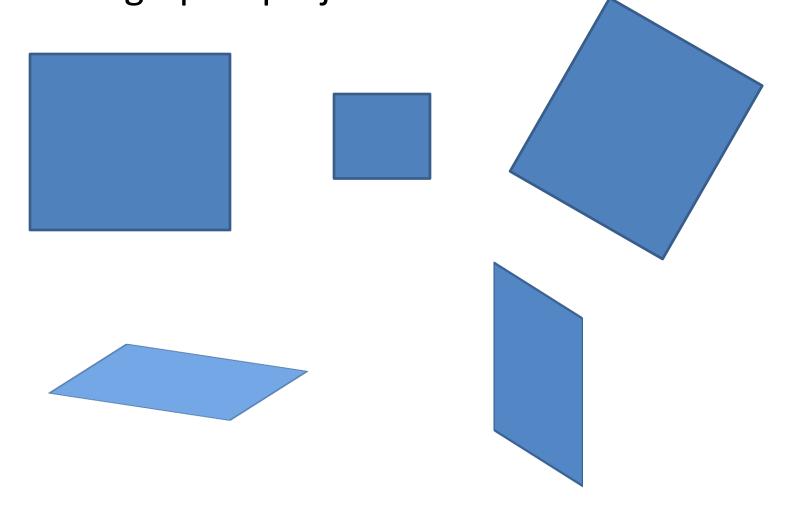


Stored Image

- 1. Match interest points from input image to database image
- Matched points vote for rough position/orientation/scale of object
- 3. Find position/orientation/scales that have at least three votes
- 4. Compute affine registration and matches using iterative least squares with outlier check
- 5. Report object if there are at least T matched points

Affine Object Model

Accounts for 3D rotation of a surface under orthographic projection



Affine Object Model

Accounts for 3D rotation of a surface under orthographic projection

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \end{bmatrix} \begin{bmatrix} x \\ y \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} x_1 & y_1 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & x_1 & y_1 & 1 \\ x_2 & y_2 & 1 & 0 & 0 & 0 \\ & \vdots & & & f \end{bmatrix} \begin{bmatrix} a \\ b \\ c \\ d \\ e \end{bmatrix} = \begin{bmatrix} x_1' \\ y_1' \\ x_2' \\ \vdots \end{bmatrix}$$

$$\mathbf{x} = [\mathbf{A^T A}]^{-1} \mathbf{A^T b}$$

What is the minimum number of matched points that we need?

Finding the objects (SIFT, Lowe 2004)

- 1. Match interest points from input image to database image
- 2. Get location/scale/orientation using Hough voting
 - In training, each point has known position/scale/orientation wrt whole object
 - Matched points vote for the position, scale, and orientation of the entire object
 - Bins for x, y, scale, orientation
 - Wide bins (0.25 object length in position, 2x scale, 30 degrees orientation)
 - Vote for two closest bin centers in each direction (16 votes total)
- 3. Geometric verification
 - For each bin with at least 3 keypoints
 - Iterate between least squares fit and checking for inliers and outliers
- 4. Report object if > T inliers (T is typically 3, can be computed to match some probabilistic threshold)

Examples of recognized objects





