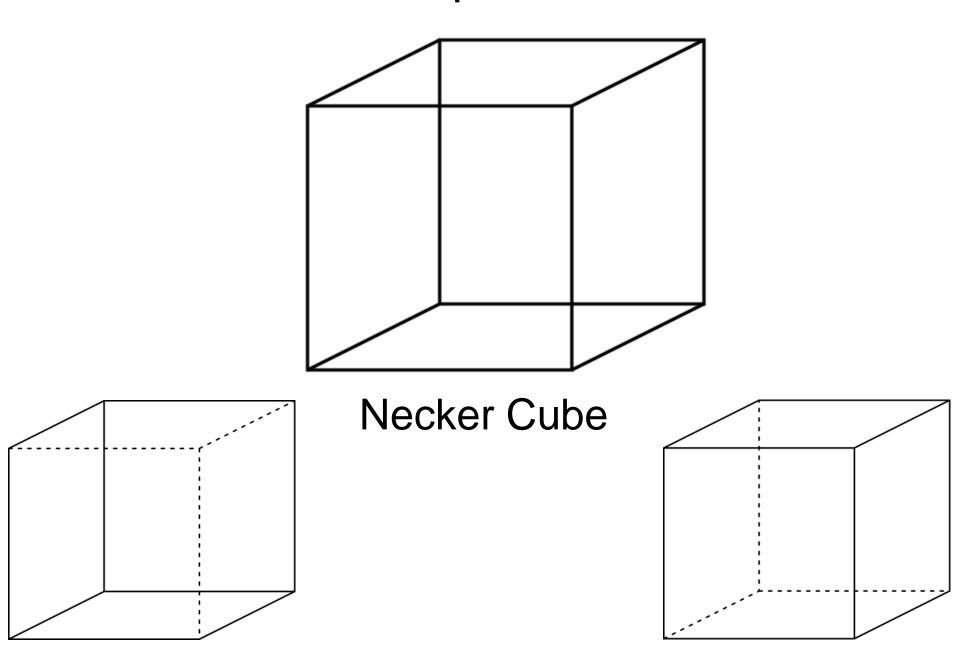
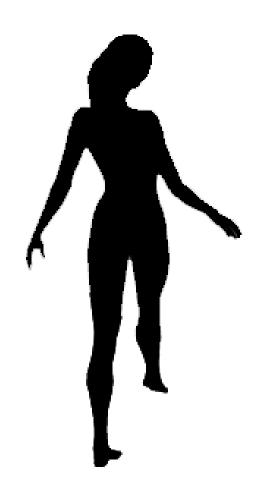
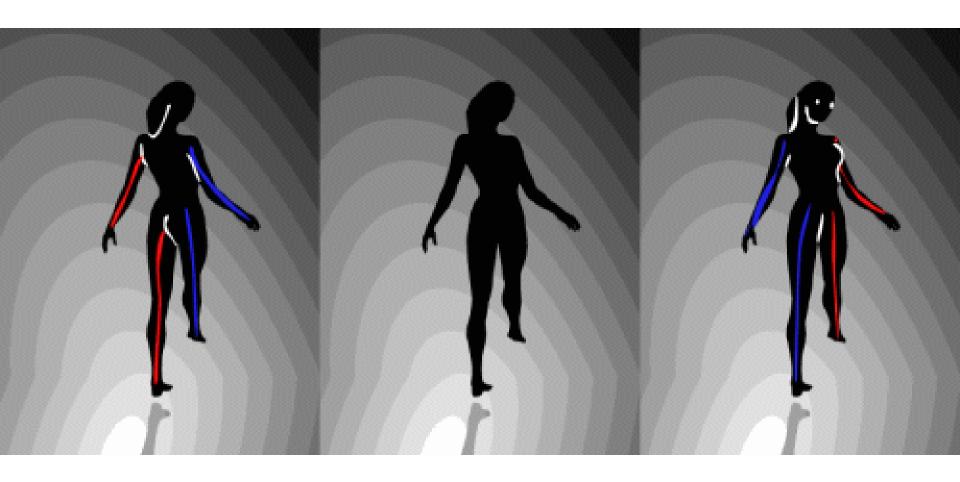
# Multi-stable Perception







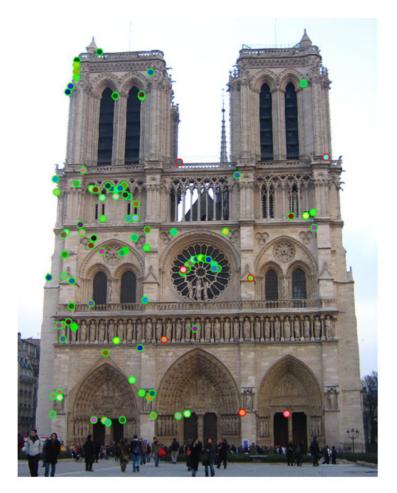
# Fitting and Alignment

Szeliski 6.1

**Computer Vision** 

James Hays

# Project 2 – due Monday





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

## 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

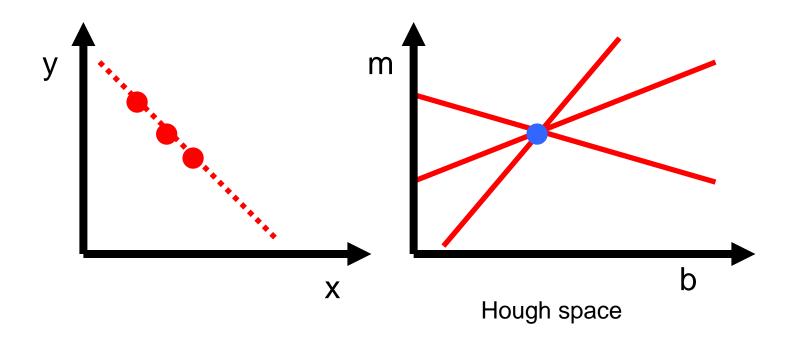
2. Each point votes for a set of parameters, incrementing those values in grid

3. Find maximum or local maxima in grid

## Review: Hough transform

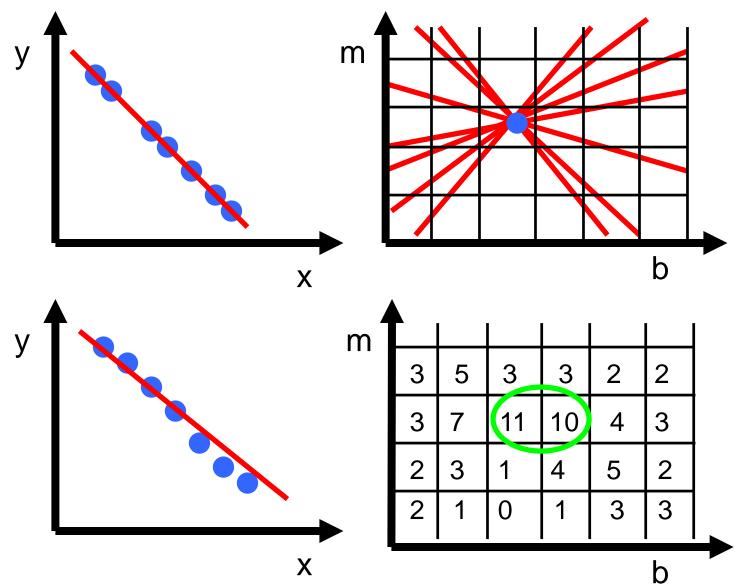
P.V.C. Hough, *Machine Analysis of Bubble Chamber Pictures,* Proc. Int. Conf. High Energy Accelerators and Instrumentation, 1959

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



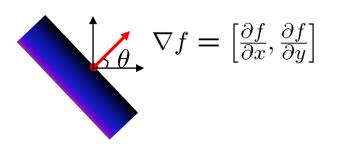
$$y = m x + b$$

# **Review: Hough transform**



# Incorporating image gradients

- Recall: when we detect an edge point, we also know its gradient direction
- But this means that the line is uniquely determined!
- Modified Hough transform:
- For each edge point (x,y)
   θ = gradient orientation at (x,y)
   ρ = x cos θ + y sin θ
   H(θ, ρ) = H(θ, ρ) + 1
   end



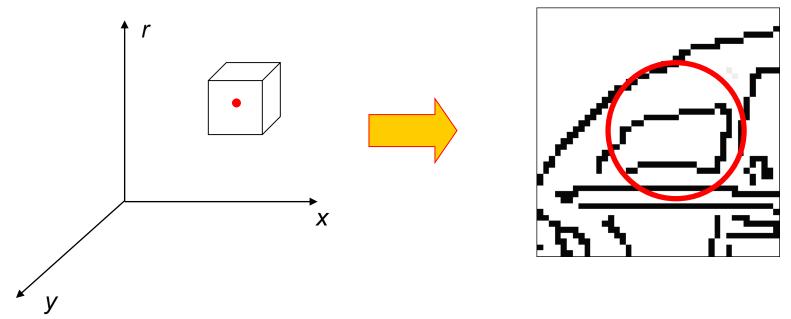
$$\theta = \tan^{-1}\left(\frac{\partial f}{\partial y} / \frac{\partial f}{\partial x}\right)$$

## Hough Transform

- How would we find circles?
  - Of fixed radius
  - Of unknown radius
  - Of unknown radius but with known edge orientation

# Hough transform for circles

 Conceptually equivalent procedure: for each (x,y,r), draw the corresponding circle in the image and compute its "support"



Is this more or less efficient than voting with features?

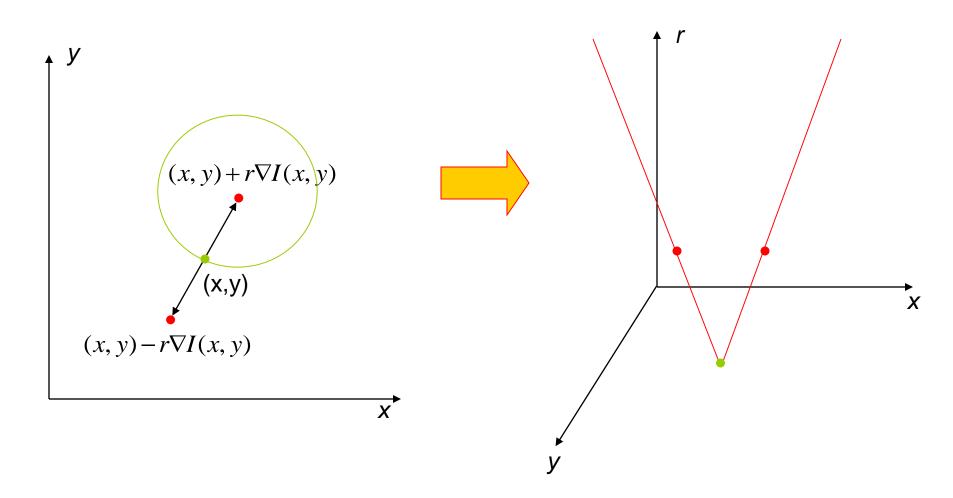
## Hough Transform

- How would we find circles?
  - Of fixed radius
  - Of unknown radius
  - Of unknown radius but with known edge orientation

# Hough transform for circles

image space

Hough parameter space



# 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

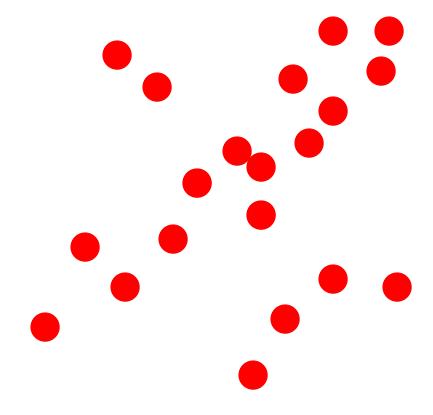
- Some sensitivity to noise
- Bin size trades off between noise tolerance, precision, and speed/memory
  - Can be hard to find sweet spot
- Not suitable for more than a few parameters
  - grid size grows exponentially

### 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):

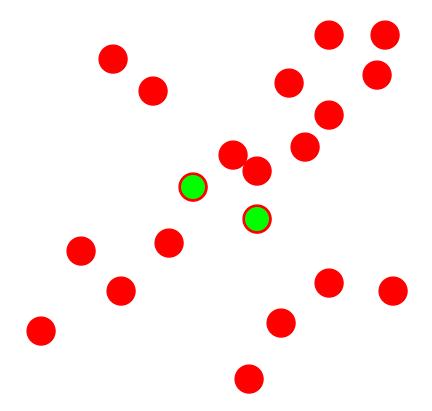
Fischler & Bolles in '81.



### 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

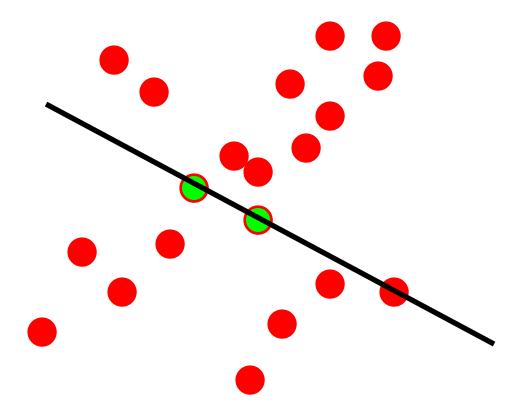
Line fitting example



### 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

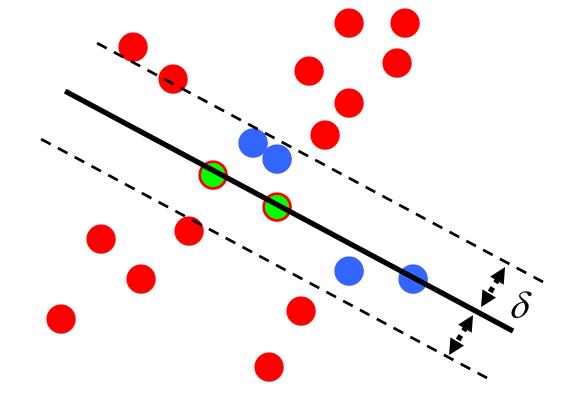
Line fitting example



### Algorithm:

- 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

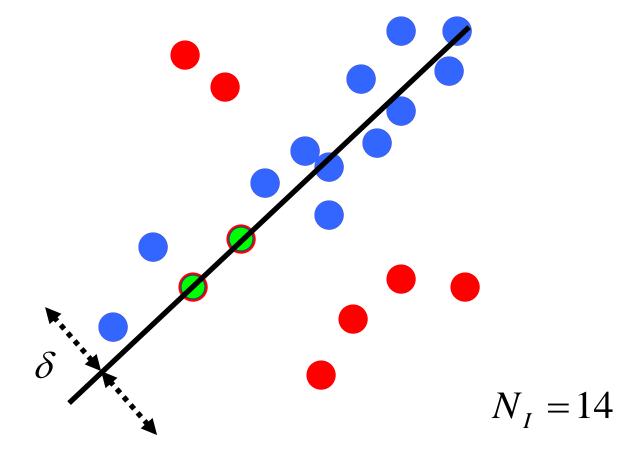
Line fitting example



$$N_{I} = 6$$

## Algorithm:

- 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



### 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

# 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  $\delta$ 
  - Choose  $\delta$  so that a good point with noise is likely (e.g., prob=0.95) within threshold
  - Zero-mean Gaussian noise with std. dev.  $\sigma$ :  $t^2=3.84\sigma^2$

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

		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		

For 
$$p = 0.99$$

## RANSAC conclusions

### Good

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

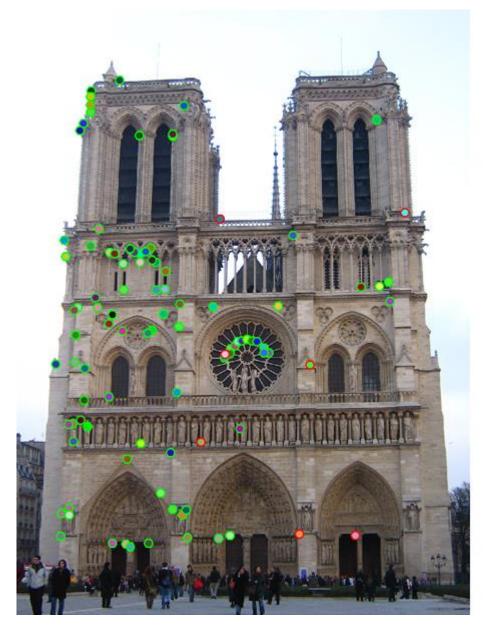
### Bad

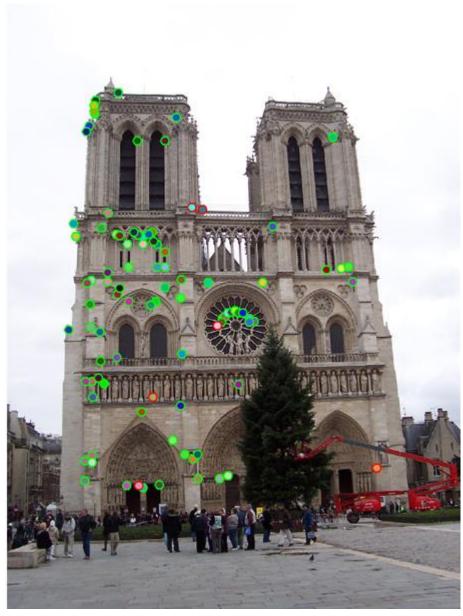
- Computational time grows quickly with fraction of outliers and number of parameters
- Not good for getting multiple fits

## Common applications

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

# How do we fit the best alignment?





# Alignment

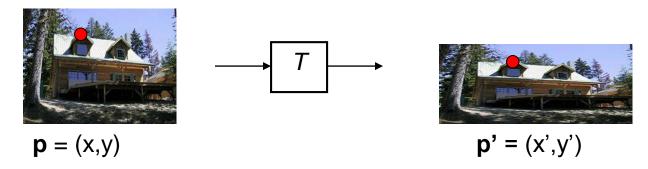
 Alignment: find parameters of model that maps one set of points to another

 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



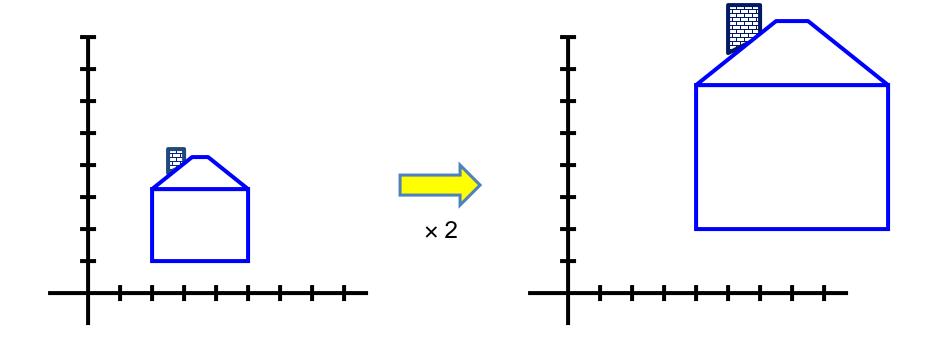
affine



perspective

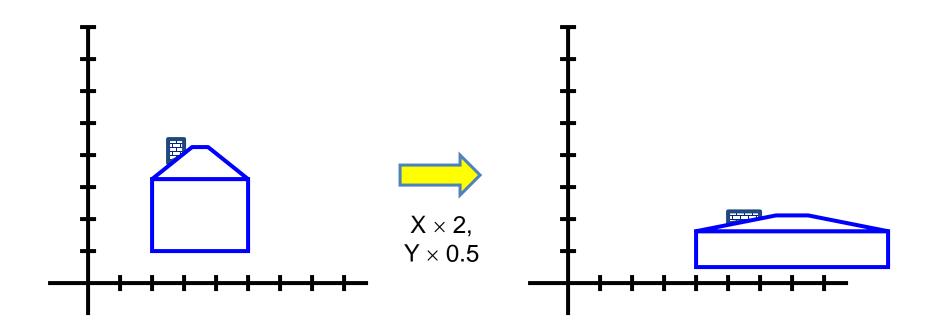
# Scaling

- 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:

$$x' = ax$$

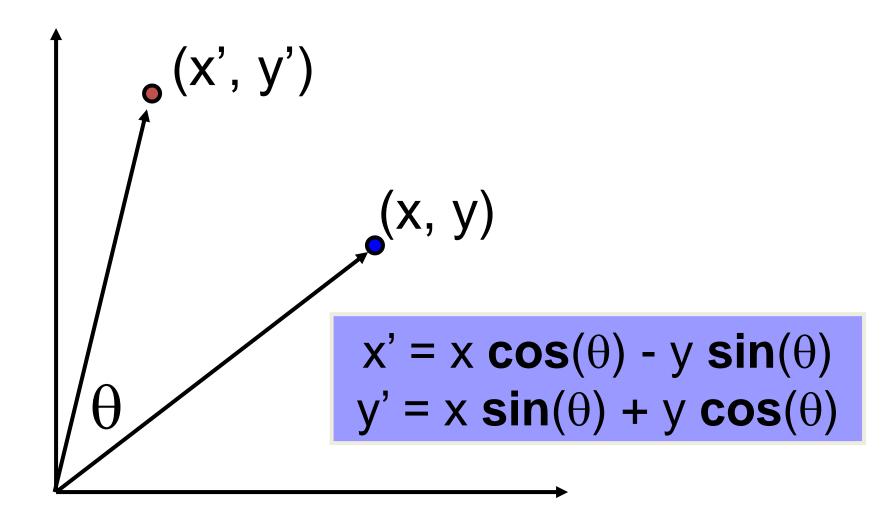
$$y' = by$$

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

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  $\theta$ ,

- -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

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

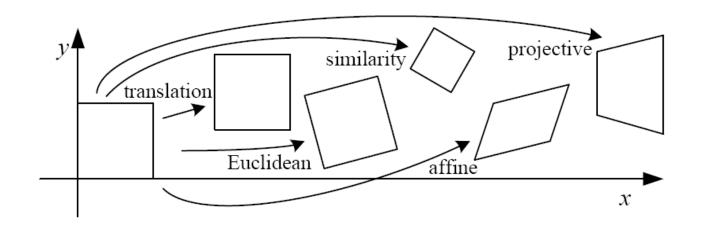
- Affine transformations, and
- Projective warps

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

#### 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)

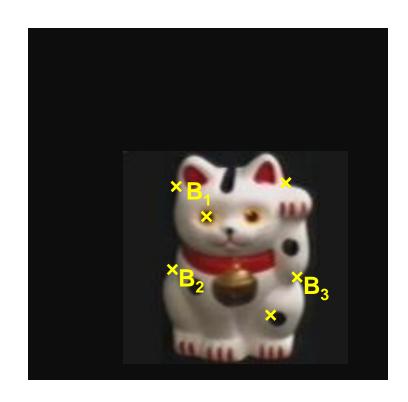
## 2D image transformations (reference table)



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

### Szeliski 2.1



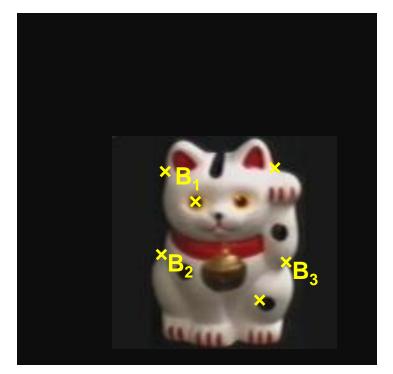


Given matched points in {A} and {B}, estimate the translation of the object

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





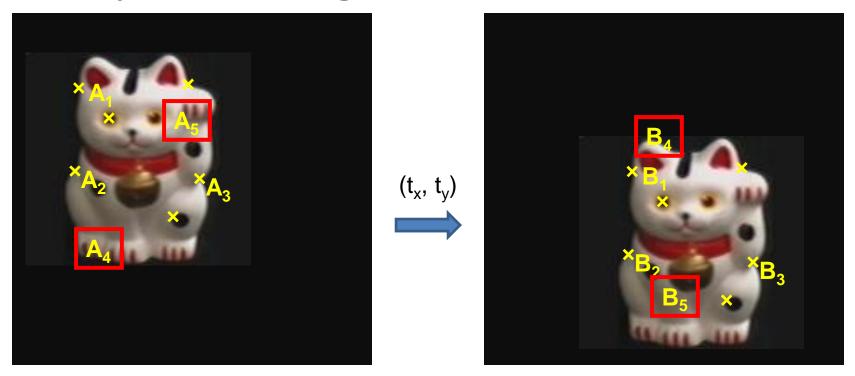


### **Least squares solution**

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

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

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

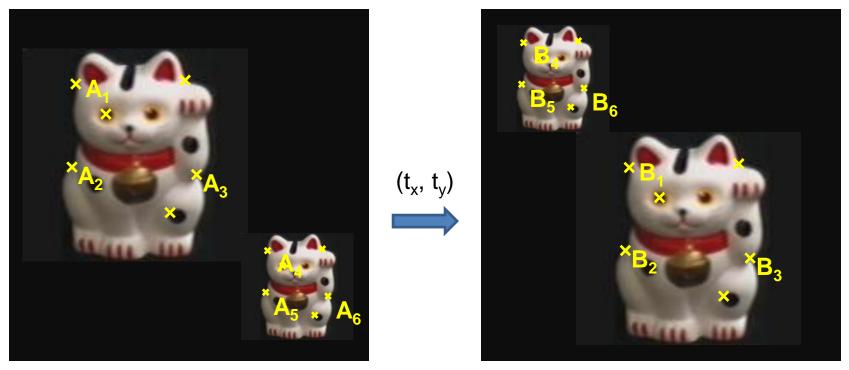


**Problem: outliers** 

#### **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

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

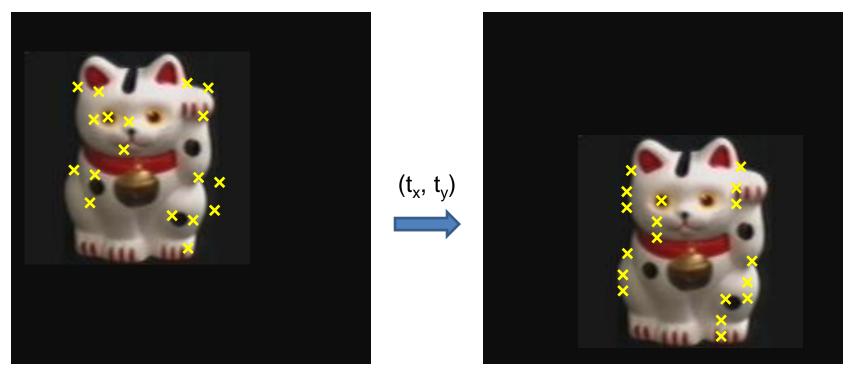


Problem: outliers, multiple objects, and/or many-to-one matches

### Hough transform solution

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

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



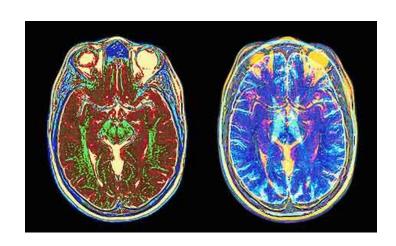
Problem: no initial guesses for correspondence

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

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