

# **Image Processing & Vision**

Lecture 06: Model Fitting – RANSAC

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#### Re-cap: Local Feature

- Keypoint is an image location at which a descriptor is computed
- Locally distinct points
- Easily localizable and identifiable
- The feature descriptor summarizes the local structure around the keypoint
- Allows us to unique matching of keypoints in presence of object pose variations, image and photometric deformations



Locally non-distinct

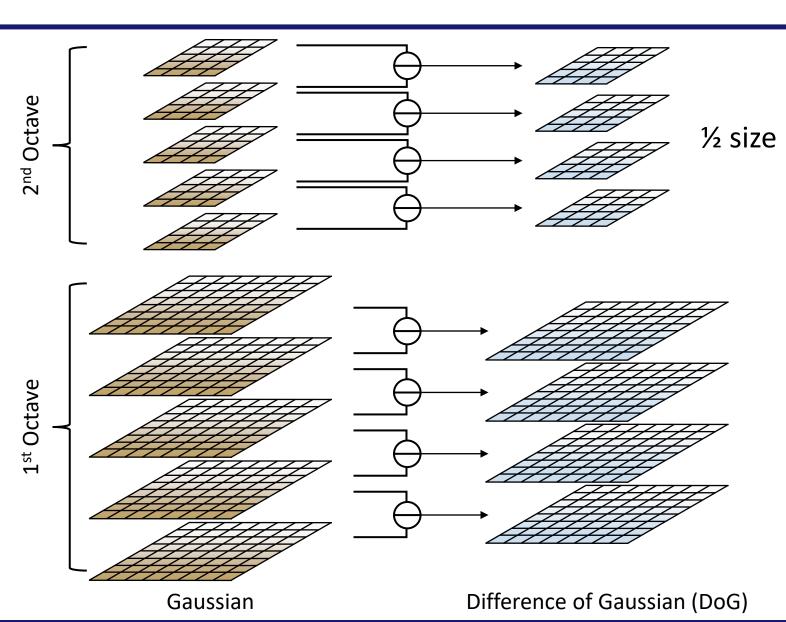
- The scale-invariant feature transform (SIFT) is an algorithm to detect, describe, and match local features in images
- SIFT describes both a detector and descriptor
- Applications: object recognition, robotic mapping and navigation, image stitching, 3D modeling, gesture recognition, video tracking, etc.
  - 1 Multi-scale extrema detection
  - ② Keypoint localization
  - 3 Orientation assignment
  - 4 Keypoint descriptor







- Begin by detecting points of interest (i.e., keypoints)
- The image is convolved with Gaussian filters at different scales
- Then the difference of successive Gaussianblurred images are taken



- Begin by detecting points of interest (i.e., keypoints)
- The image is convolved with Gaussian filters at different scales
- Then the difference of successive Gaussianblurred images are taken
- Keypoints are taken as maxima/minima of the DoG at multiple scales

Gaussian variance The largest one is selected Scale of Difference of Gaussian (DoG)

- Scale-space extrema detection produces too many keypoint candidates, some of which are unstable
- In keypoint localization, we reject points which are low contrast (and are therefore sensitive to noise) or poorly localized along an edge

- How do we decide whether a keypoint is poorly localized or well-localized?
- In SIFT, compute the ratio of the eigenvalues of covariance matrix and

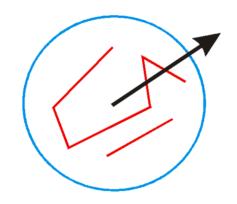
check if it is greater than a threshold

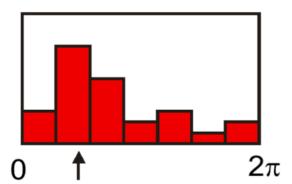


- Each keypoint is assigned one or more orientations based on local image gradient directions
- This is the key step in achieving invariance to rotation on the Gaussian-smoothed image,  $\boldsymbol{L}$

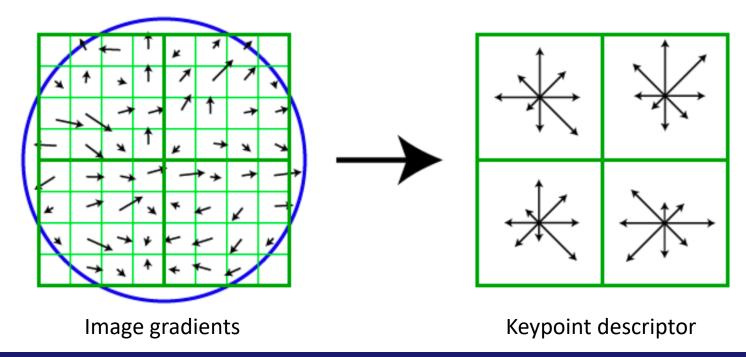
$$m(x,y) = \sqrt{(L(x+1,y) - L(x-1,y))^2 + (L(x,y+1) - L(x,y-1))^2}$$

$$\theta(x,y) = \tan^{-1} \left[ \left( L(x+1,y) - L(x-1,y) \right) / \left( L(x,y+1) - L(x,y-1) \right) \right]$$





- Thresholded image gradients are sampled over 16 x 16 array of locations in scale space (weighted by a Gaussian with sigma half the size of the window)
- Create array of orientation histograms
- 8 Orientations x (4 x 4) histogram array



#### 1 Scale-space representation and local extreme detection

- Use DoG/LoG Pyramid
- 3 scales/octave, down-sample by factor of 2 each octave

#### ② Keypoint localization

 Select stable keypoints by thresholding on the magnitude of extrema and ratio of principal curvatures

#### **③ Keypoint orientation assignment**

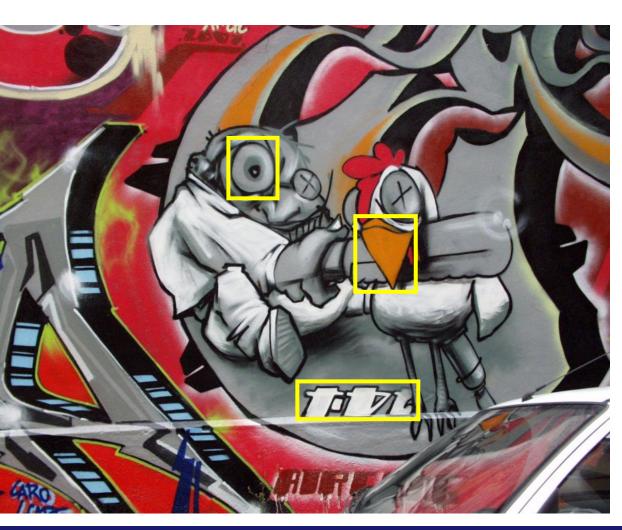
Based on histogram of local image gradient directions

#### 4 Keypoint descriptor

- Histogram of local gradient directions Vector with  $8 \times (4 \times 4) = 128$  dimensions
- Vector normalization

# Re-cap: Application of Good Local Features

Feature Matching





#### Re-cap: Application of Good Local Features

- Example of Feature Matching
  - Composed of more than 1,000 images and carefully assembled over the ensuing months



NASA's Curiosity rover captured its highest-resolution panorama of the Martian surface

## Re-cap: Application of Good Local Features

Object Tracking & Recognition



# **Topics**

Model Fitting – RANSAC

<sup>\*</sup>Note: Many of these slides in this course were adapted from Computer Vision at CMU (16-385) and UBC (CPSC425)

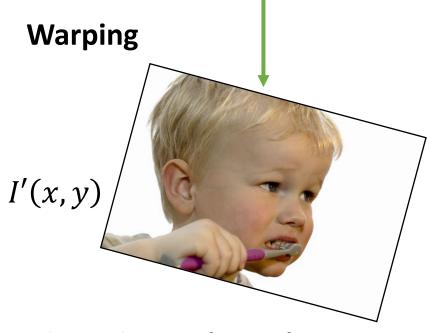
# Transformation: Warping

- The model (warping) gives us a way to transform any pixel in the original image I(x,y) to the corresponding image I'(x,y)
- We will call this "Warping" a "Model"

$$\begin{bmatrix} X' \\ Y' \\ 1 \end{bmatrix} = M \begin{bmatrix} X \\ Y \\ 1 \end{bmatrix}$$

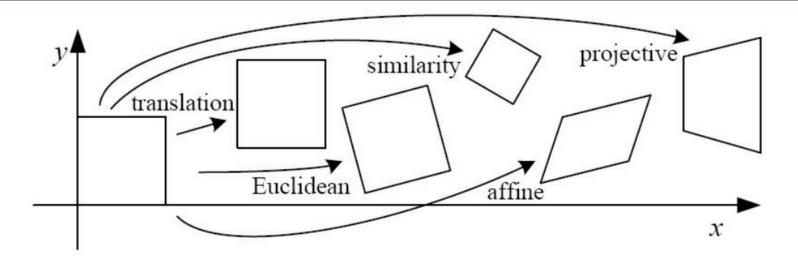
$$I(x,y) = I'(x',y')$$





Change domain of image function

# Forms of the Model (Warping)

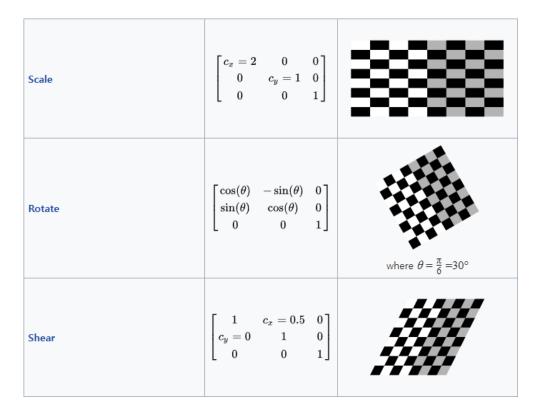


Name	Matrix	# D.O.F.	Preserves:	Icon
translation	$egin{bmatrix} I & I & I \end{bmatrix}_{2 imes 3}$			
rigid (Euclidean)	$egin{bmatrix} egin{bmatrix} oldsymbol{R} oldsymbol{t} \end{bmatrix}_{2 imes 3}$		_	$\Diamond$
similarity	$\begin{bmatrix} sR \mid t \end{bmatrix}_{2 \times 3}$	_	_	$\Diamond$
affine	$\left[egin{array}{c} oldsymbol{A} \end{array} ight]_{2 imes 3}$			
projective	$\left[egin{array}{c}  ilde{H} \end{array} ight]_{3 imes 3}$			

#### **Affine Model**

- In Euclidean geometry, an affine transformation is a geometric transformation that preserves lines and parallelism
- But not necessarily distances and angles

Identity (transform to original image)	$\begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}$	
Translation	$egin{bmatrix} 1 & 0 & v_x > 0 \ 0 & 1 & v_y = 0 \ 0 & 0 & 1 \end{bmatrix}$	338
Reflection	$egin{bmatrix} -1 & 0 & 0 \ 0 & 1 & 0 \ 0 & 0 & 1 \end{bmatrix}$	



#### **Affine Model**

• Affine transform of [x, y] to [u, v]:

$$\begin{bmatrix} u \\ v \end{bmatrix} = \begin{bmatrix} m_1 & m_2 \\ m_3 & m_4 \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix} + \begin{bmatrix} t_x \\ t_y \end{bmatrix}$$

Rewrite to solve the transformation parameters:

$$\begin{bmatrix} x_1 & y_1 & 0 & 0 & 1 & 0 \\ 0 & 0 & x_1 & y_1 & 0 & 1 \\ x_2 & y_2 & 0 & 0 & 1 & 0 \\ 0 & 0 & x_2 & y_2 & 0 & 1 \\ & & \cdots & \cdots & & \\ & & & t_y \end{bmatrix} \begin{bmatrix} m_1 \\ m_2 \\ m_3 \\ m_4 \\ t_x \\ t_y \end{bmatrix} = \begin{bmatrix} u_1 \\ v_1 \\ u_2 \\ v_2 \\ \cdots \\ \cdots \end{bmatrix}$$

6 equations, 6 unknowns

#### **Affine Model**

• Suppose we have  $k \geq 3$  matches,  $[x_i, y_i]$  to  $[u_i, v_i]$  where  $i = 1, 2, \dots, k$ , then,

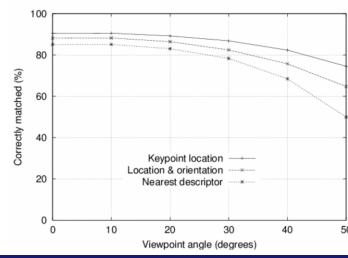
$$\begin{bmatrix} x_1 & y_1 & 0 & 0 & 1 & 0 \\ 0 & 0 & x_1 & y_1 & 0 & 1 \\ x_2 & y_2 & 0 & 0 & 1 & 0 \\ 0 & 0 & x_2 & y_2 & 0 & 1 \\ & & \cdots & \cdots & & \\ x_k & y_k & 0 & 0 & 1 & 0 \\ 0 & 0 & x_k & y_k & 0 & 1 \end{bmatrix} \begin{bmatrix} m_1 \\ m_2 \\ m_3 \\ m_4 \\ t_x \\ t_y \end{bmatrix} = \begin{bmatrix} u_1 \\ v_1 \\ u_2 \\ v_2 \\ \cdots \\ u_k \\ v_k \end{bmatrix}$$

#### Limitation

- We need to have exact matches
- Only 3 keypoints are needed for recognition, so extra keypoints provide robustness
- It is very **difficult** to find exact match
- If we can find exact match **80**% of the time, we can find 3 matches

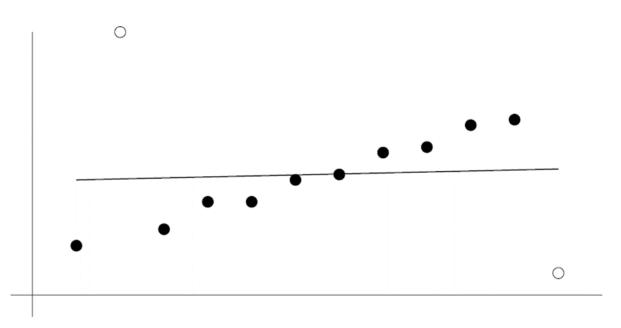
correctly only about 50% of the time.

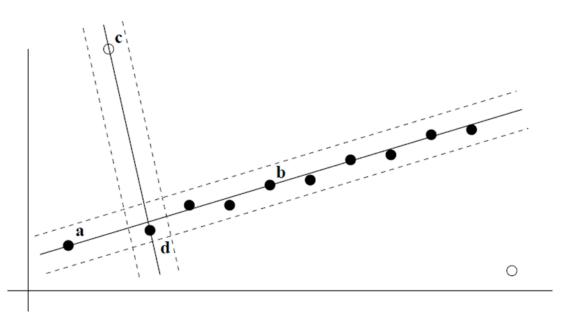
- Image noise, deformations, will make this worse
- Multiple object instances will make this impossible



#### Fitting a Model to Noisy Data

- Problem of a Least-Square Line Fitting
- In a line fit to the data
- In a classification of the data into inliers (valid points) and outliers





A least-squares line fitting

**RANSAC** line fitting

#### Fitting a Model to Noisy Data

- Suppose we are fitting a line to a dataset that consists of 50% outliers.
- We can fit a line using two points

If we draw pairs of points uniformly at random, what fraction of pairs will consist entirely of **good data points (inliers)?** 

#### Fitting a Model to Noisy Data

- Suppose we are fitting a line to a dataset that consists of 50% outliers.
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If we draw pairs of points uniformly at random, what fraction of pairs will consist entirely of **good data points (inliers)?** 

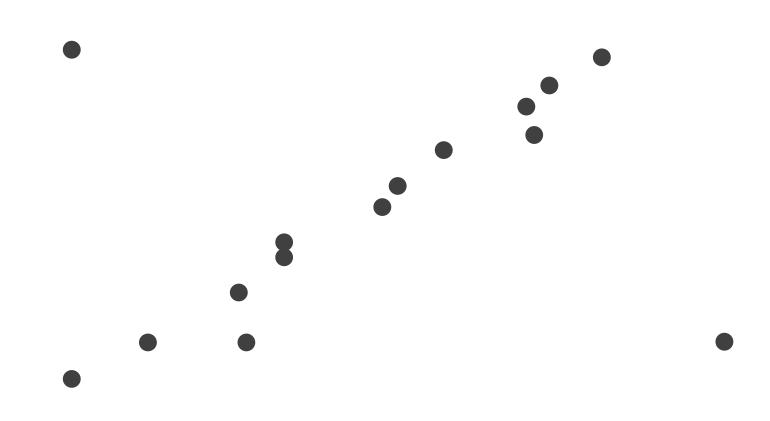
- If we draw pairs of points uniformly at random, then about 1/4 of these pairs will consist entirely of 'good' data points (inliers)
- We can identify these good pairs by noticing that a large collection of other points lie close to the line fitted to the pair
- A better estimate of the line can be obtained by refitting the line to the points that lie close to the line

Objective: Robust fit of a model to a data set S which contains outliers

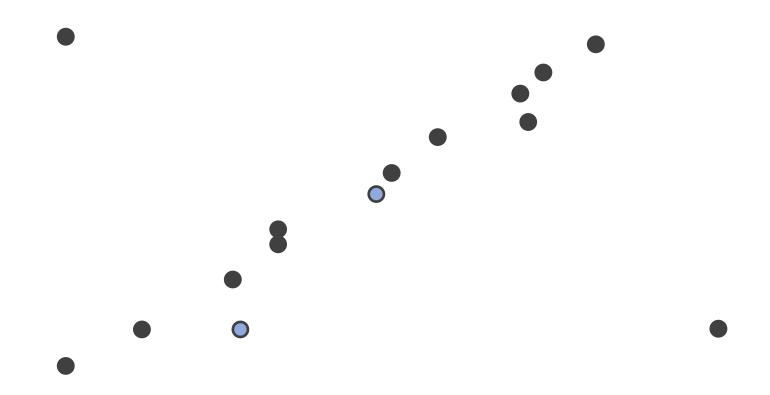
#### Algorithm:

- Randomly choose minimal subset of data points necessary to fit model (a sample)
- ② Points within some distance threshold, t, of model are a **consensus set**. Size of consensus set is model's **support**
- $\odot$  Repeat for N samples; model with biggest support is most robust fit.
  - Points within distance t of best model are inliers
  - Fit final model to all inliers

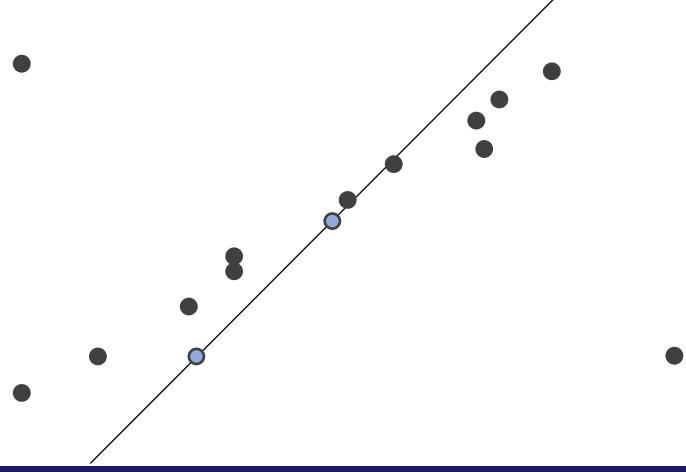
RANSAC Line Fitting Example



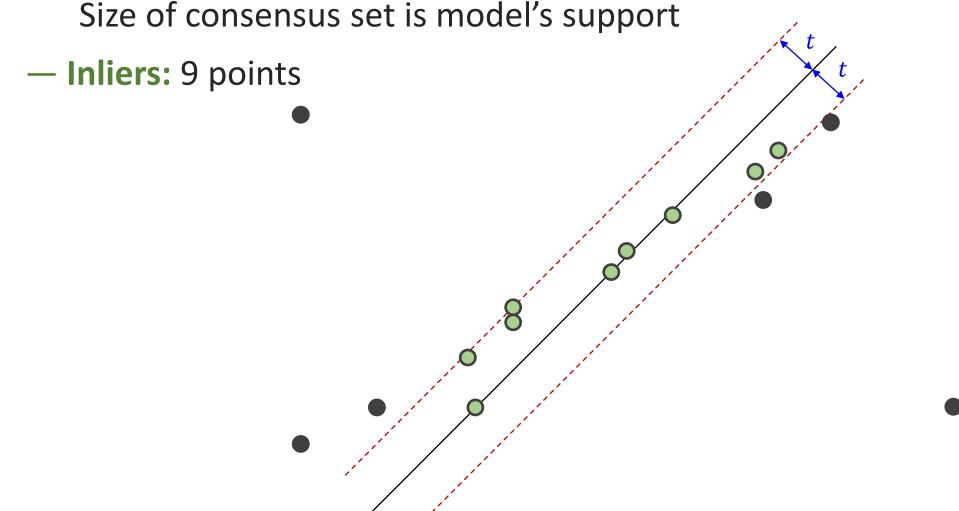
1 Randomly choose minimal subset of data points necessary to fit model (a sample)



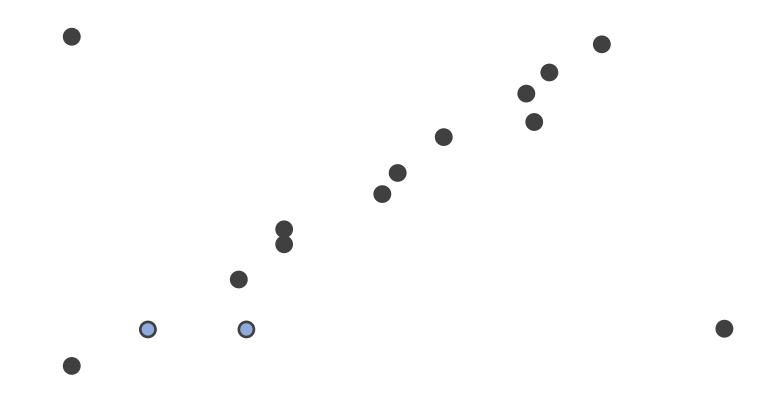
1 Randomly choose minimal subset of data points necessary to fit model (a sample)



② Points within some distance threshold, t, of model are a consensus set.

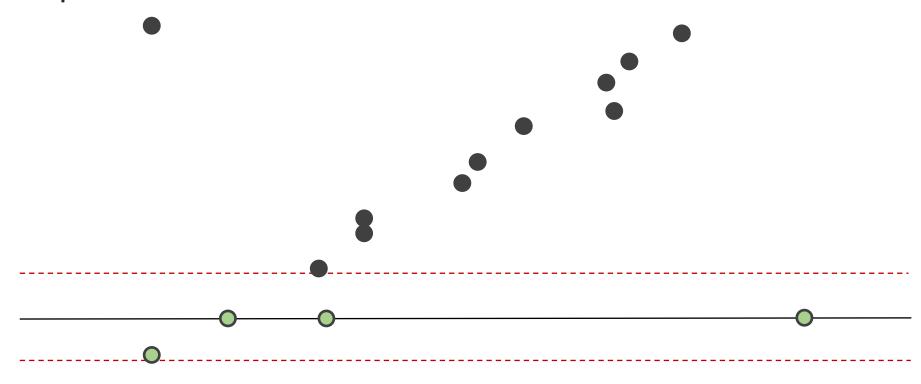


1 Randomly choose minimal subset of data points necessary to fit model (a sample)

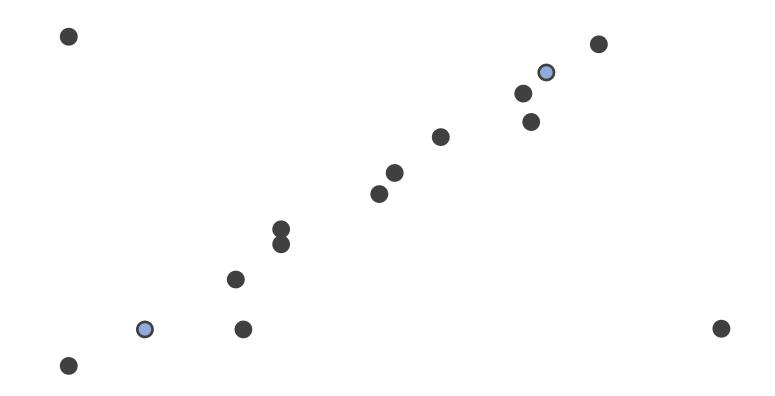


② Points within some distance threshold, t, of model are a consensus set. Size of consensus set is model's support

— Inliers: 4 points



1 Randomly choose minimal subset of data points necessary to fit model (a sample)



② Points within some distance threshold, t, of model are a consensus set.

Size of consensus set is model's support — Inliers: 12 points

Repeat for N samples; model with biggest support is most robust fit Points within distance t of best model are inliers Fit final model to all inliers • : Inliers : Outliers

### **How Many Samples?**

- Let  $\omega$  be the fraction of inliers (i.e., points on line)
- Let n be the number of points needed to define hypothesis (n=2 for a line in the plane)
- Suppose k samples are chosen

- The probability that a single sample of n points is correct (all inliers) is  $\omega^n$
- The probability that all k samples fail is  $(1 \omega^n)^k$

Choose k large enough to keep this below a target failure rate

# RANSAC: k Samples Chosen (p = 0.99)

Sample size	Proportion of outliers							
n	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	

#### **After RANSAC**

 RANSAC divides data into inliers and outliers and yields estimate computed from minimal set of inliers

- Improve this initial estimate with estimation over all inliers (e.g., with standard least-squares minimization)
- But this may change inliers, so alternate fitting with re-classification as inlier/outlier

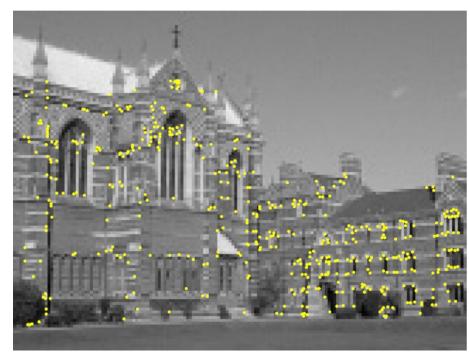
- How to get correct correspondences without human intervention?
- RANSAC can be used for image stitching or automatic determination of epipolar geometry





Left image Right image

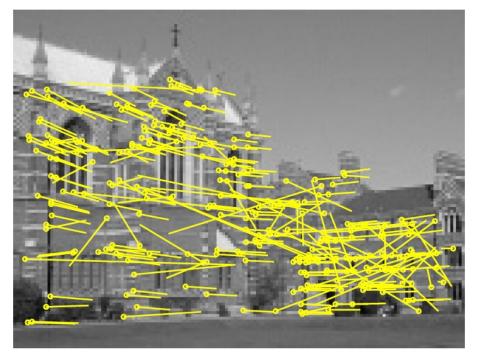
- (1) Feature Extraction
- Find features in pair of images using Harris corner detector
- Assumes images are roughly the same scale





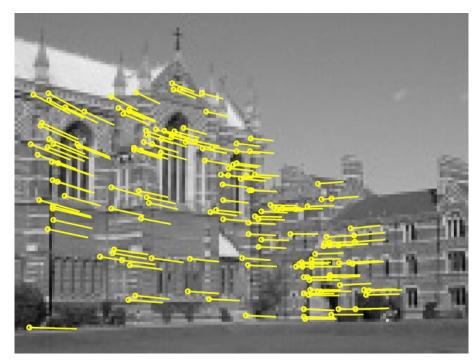
Left image Right image

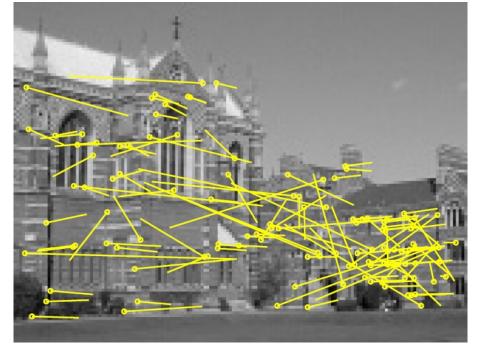
- 2 Finding Feature Matches Initial Match Hypothesis
- 268 matched features over SSD threshold superimposed on left image (pointing to locations of corresponding feature in right image)



268 matched features on left image

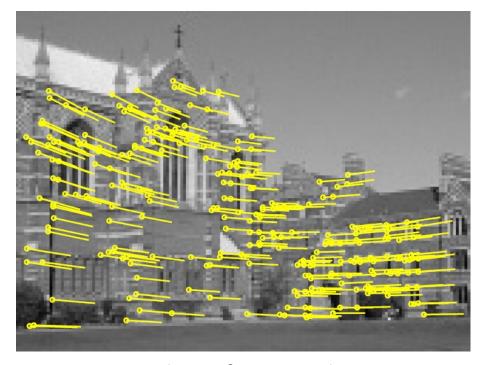
- (3) Outliers & Inliers after RANSAC
- -n is 4 for this problem (a homography relating 2 images)
- 43 samples used with t = 1.25 pixels (Assume up to 45% outliers)





151 inliers 117 outliers

- 4 Final Matches
- Maximum Likelihood Estimation (MLE)
- Guided Matching



Final set of 262 matches

#### **Summary:** RANSAC

#### Advantages:

- General method suited for a wide range of model fitting problems
- Easy to implement and easy to calculate its failure rate

#### Disadvantages:

- Only handles a moderate percentage of outliers without cost blowing up
- Many real problems have high rate of outliers (but sometimes selective choice of random subsets can help)