

Chapter 06 Robust Estimation of Homography

Lin ZHANG
School of Computer Science and Technology
Tongji University



- Problem of Homography Estimation
- RANSAC-based Homography Estimation



Problem definition:

On two projective planes π_1 and π_2 , there is a set of corresponding points $\left\{\mathbf{x}_i, \mathbf{x}_i^{'}\right\}_{i=1}^n$, and we suppose that there is a homography matrix linking the two planes,

$$c_i \mathbf{x}_i' = H \mathbf{x}_i, i = 1, 2, ..., n$$

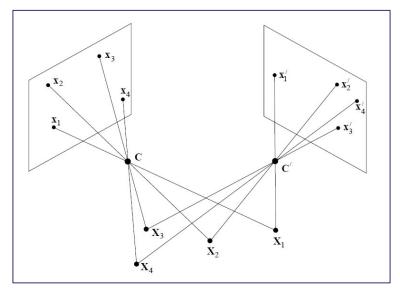
Coordinates of $\left\{\mathbf{x}_i\right\}_i^n$ and $\left\{\mathbf{x}_i^{'}\right\}_{i=1}^n$ are known, we need to find H

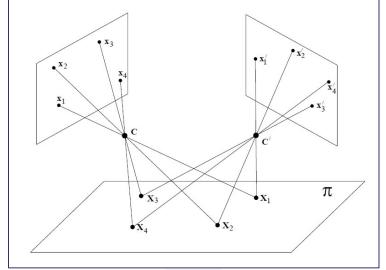
$$H = \begin{bmatrix} h_{11} & h_{12} & h_{13} \\ h_{21} & h_{22} & h_{23} \\ h_{31} & h_{32} & h_{33} \end{bmatrix}$$

Note: *H* is defined up to a scale factor. In other words, it has 8 DOFs



Note: Theoretically speaking, homography can only be estimated between two planes, i.e., when you use such a technique to stitch two images, image contents should be roughly on the same plane











4 point-correspondence pairs can uniquely determine a homography matrix since each correspondence pair solves two degrees of freedom

$$c\begin{pmatrix} x \\ y \\ 1 \end{pmatrix} = \begin{bmatrix} h_{11} & h_{12} & h_{13} \\ h_{21} & h_{22} & h_{23} \\ h_{31} & h_{32} & h_{33} \end{bmatrix} \begin{pmatrix} u \\ v \\ 1 \end{pmatrix} \longrightarrow \begin{cases} h_{11}u + h_{12}v + h_{13} = cx \\ h_{21}u + h_{22}v + h_{23} = cy \\ h_{31}u + h_{32}v + h_{33} = c \end{cases}$$

$$\begin{cases} \frac{h_{11}u + h_{12}v + h_{13}}{h_{31}u + h_{32}v + h_{33}} = x \\ \frac{h_{21}u + h_{22}v + h_{23}}{h_{31}u + h_{32}v + h_{33}} = y \end{cases}$$

Note: here we assume that the matching points are all finite points (no points at infinity)



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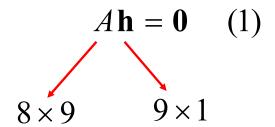
wo degrees of freedom
$$\begin{pmatrix} u & v & 1 & 0 & 0 & 0 & -ux & -vx & -x \\ 0 & 0 & 0 & u & v & 1 & -uy & -vy & -y \end{pmatrix} \begin{pmatrix} h_{11} \\ h_{12} \\ h_{21} \\ h_{22} \\ h_{23} \\ h_{31} \\ h_{32} \\ h_{33} \end{pmatrix} = \mathbf{0}$$
 Thus, four correspondence pairs generate 8 equations

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$$\begin{vmatrix} h_{12} \\ h_{13} \\ h_{21} \\ h_{22} \\ h_{23} \\ h_{31} \\ h_{32} \\ h_{33} \end{vmatrix} = \mathbf{0}$$



4 point-correspondence pairs can uniquely determine a homography matrix since each correspondence pair solves two degrees of freedom



Normally, Rank(A) = 8; thus (1) has 1 (9-8) solution vector (linear independent) in its solution space

In our case, since we have n>4 point pairs, we get

$$\mathbf{A}_{2n\times 9}\mathbf{h}_{9\times 1}=\mathbf{0}$$

It is an overdetermined homogeneous linear equation system



Since only the ratios among the elements of H take effect, in another way we can fix $h_{33}=1$ (suppose that $h_{33}!=0$),

$$c \begin{pmatrix} x \\ y \\ 1 \end{pmatrix} = \begin{bmatrix} h_{11} & h_{12} & h_{13} \\ h_{21} & h_{22} & h_{23} \\ h_{31} & h_{32} & 1 \end{bmatrix} \begin{pmatrix} u \\ v \\ 1 \end{pmatrix} \longrightarrow \begin{cases} h_{11}u + h_{12}v + h_{13} = cx \\ h_{21}u + h_{22}v + h_{23} = cy \\ h_{31}u + h_{32}v + 1 = c \end{cases} \longrightarrow \begin{cases} \frac{h_{11}u + h_{12}v + h_{13}}{h_{31}u + h_{32}v + 1} = x \\ \frac{h_{21}u + h_{22}v + h_{23}}{h_{31}u + h_{32}v + 1} = y \end{cases}$$



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- When there are more than 4 correspondence pairs, is it a proper way to use the LS method to solve the model directly?
 - ➤ NO! Because usually, outliers exist among the correspondence pairs

RANdom SAmple Consensus (RANSAC) is an iterative framework to estimate a parametric model from observations with noisy outliers



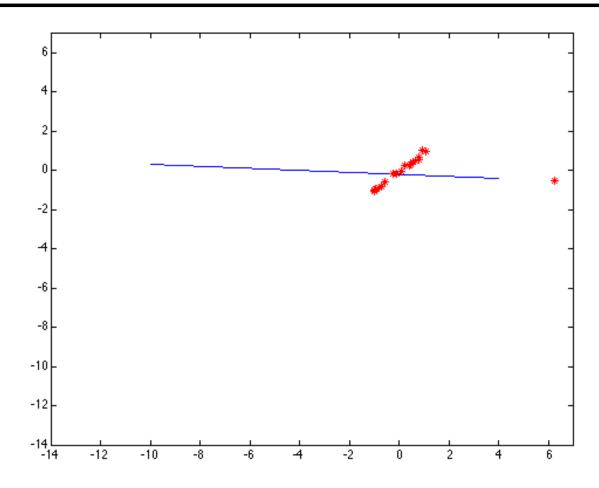
Objective

Robust fit a model to a data set S which contains outliers

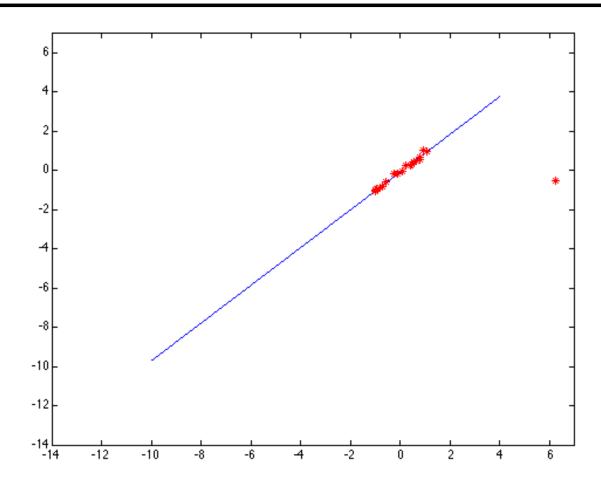
Algorithm

- (1) Randomly select a sample of s data points from S and instantiate the model from this subset
- (2) 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 the sample and defines the inliers of S
- (3) If the size of S_i (the number of inliers) is greater than some threshold T, re-estimate the model using all points in S_i and terminate
- (4) If the size of S_i is less than T, select a new subset and repeat the above
- (5) After N trials the largest consensus set S_i is selected, and the model is re-estimated using all points in the subset S_i

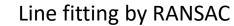


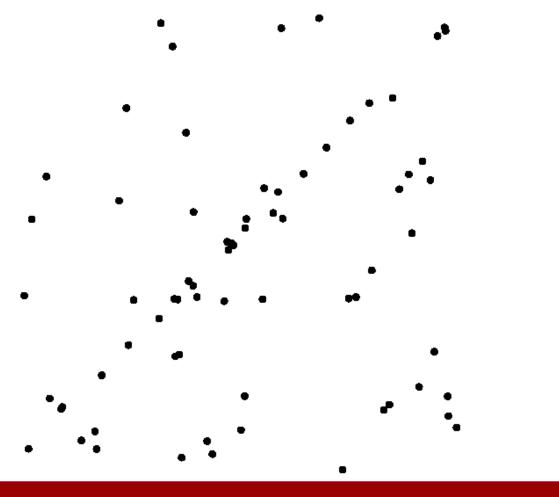






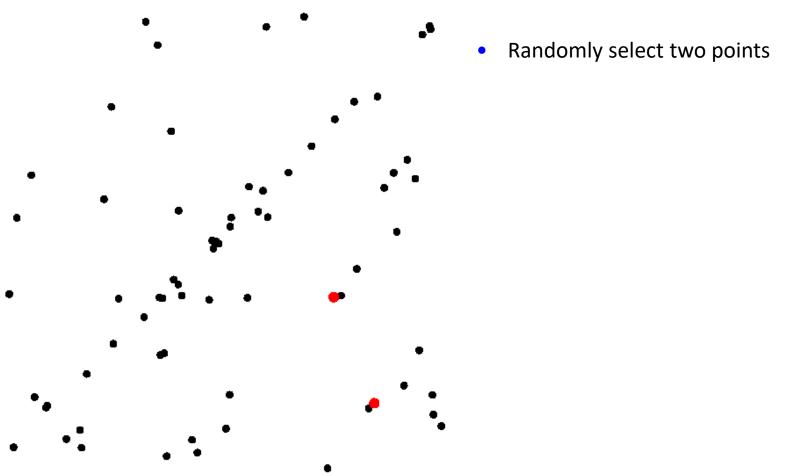




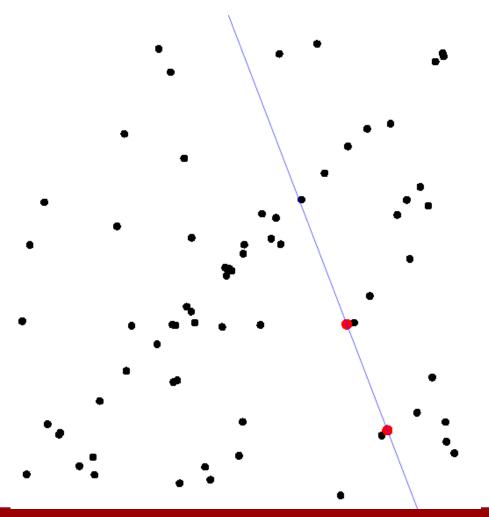








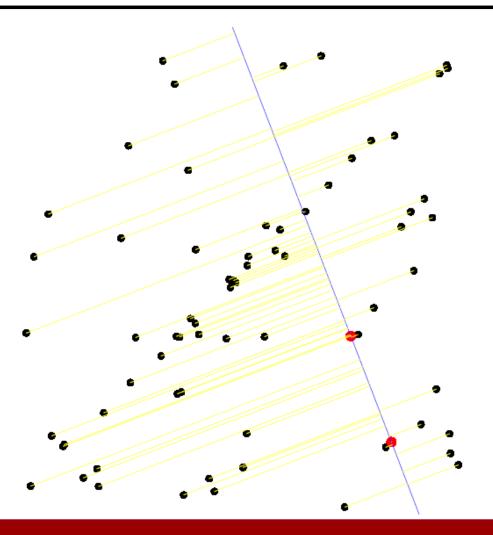




Line fitting by RANSAC

- Randomly select two points
- The hypothesized model is the line passing through the two points

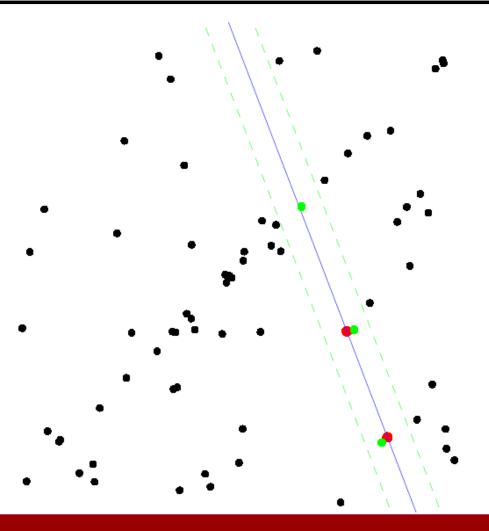




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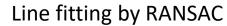


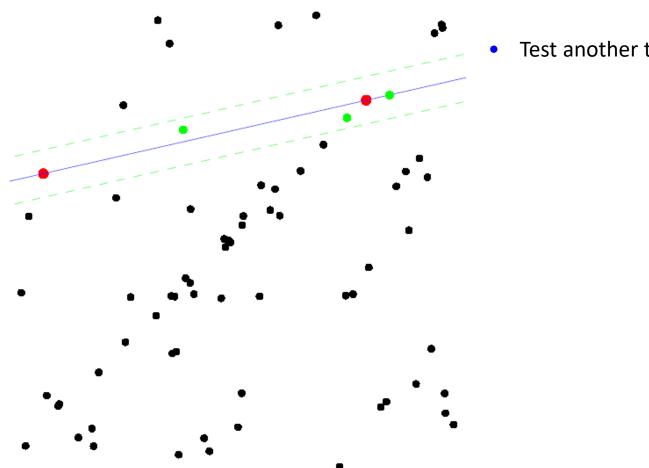


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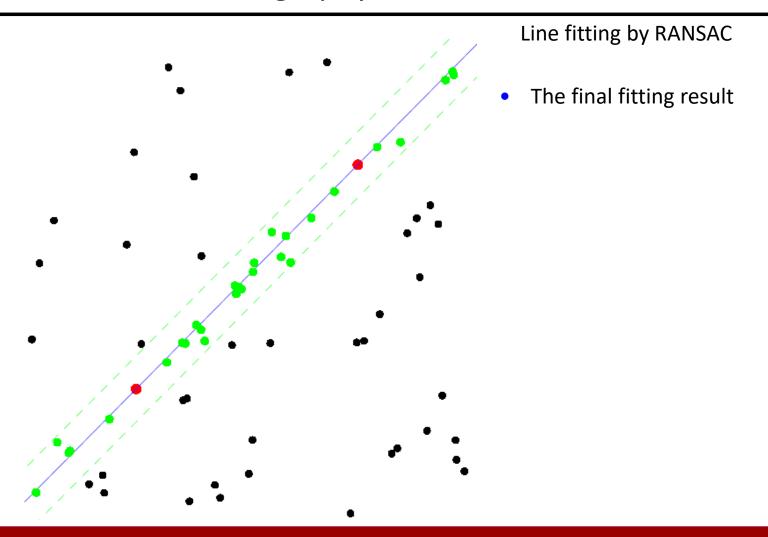




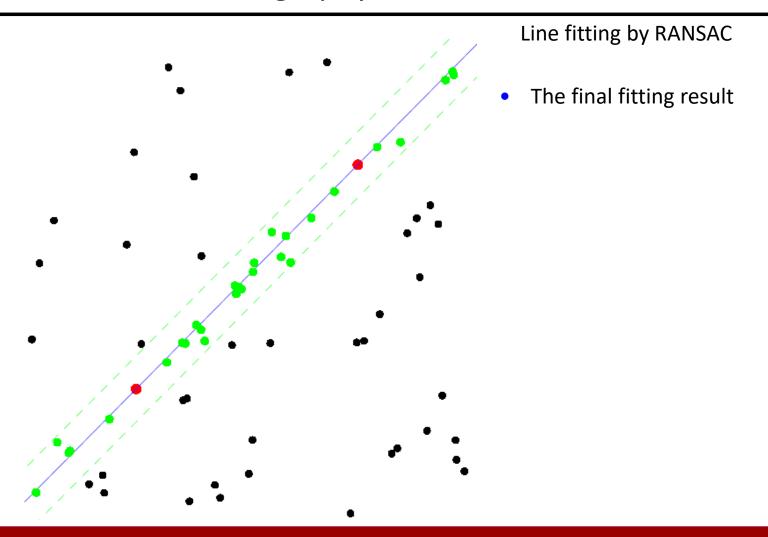


Test another two points

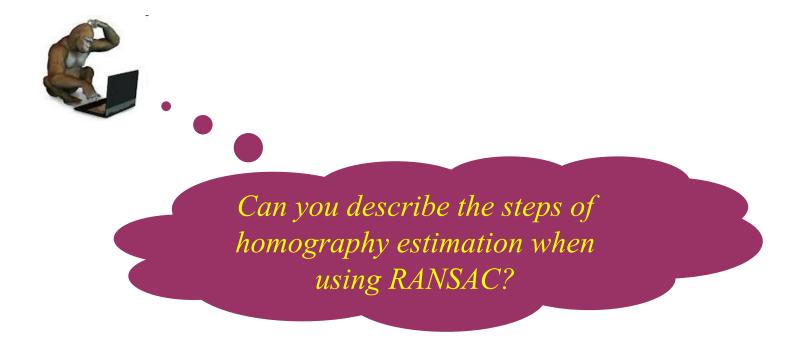










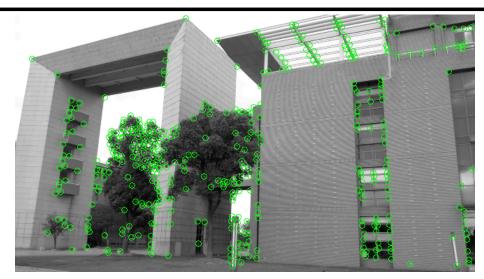




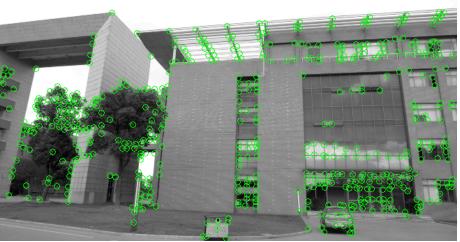








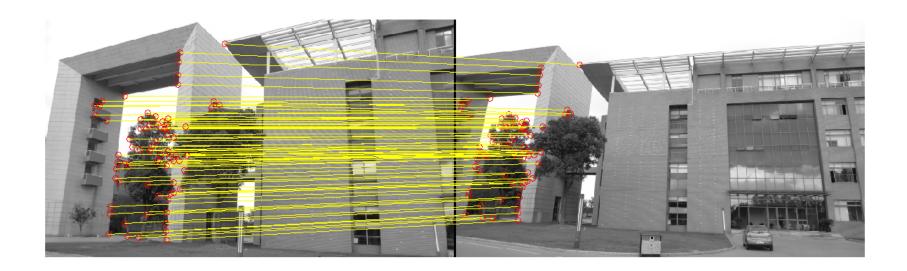
Interest points detection





Correspondence estimation

Then, the homography matrix can be estimated by using the correspondence pairs with RANSAC





Transform image one using the estimated homography matrix





Finally, stitch the transformed image one with image two





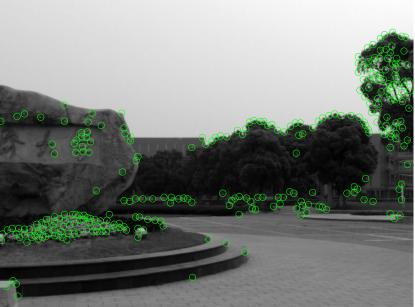








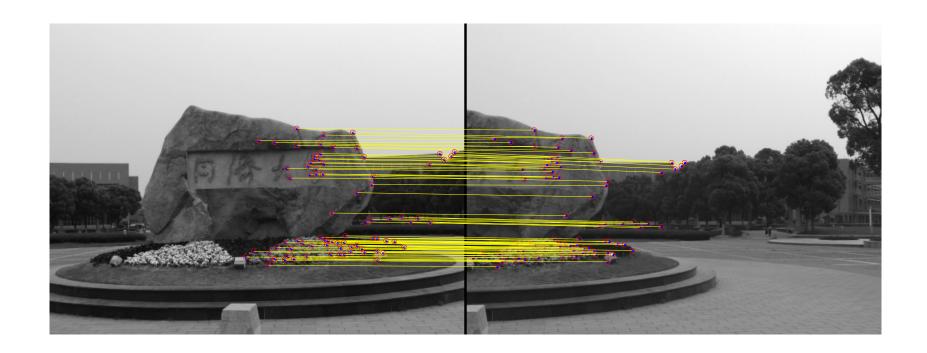
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