

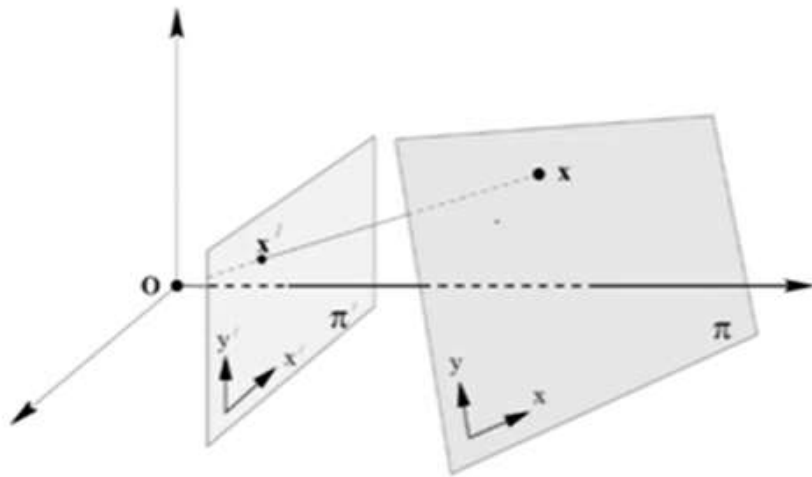
# Projective Homography

## RANSAC Algorithm

Some slides were adapted/taken from various sources, including 3D Computer Vision of Prof. Hee, NUS, Air Lab Summer School, The Robotic Institute, CMU, Computer Vision of Prof. Mubarak Shah, UCF, Computer Vision of Prof. William Hoff, Colorado School of Mines and many more. We thankfully acknowledge them. Students are requested to use this material for their study only and **NOT** to distribute it.

# Planner Projective Transformations

- **Central projection** maps points on one plane to points on another plane.
- And represented by a **linear mapping** of homogeneous coordinates  $\mathbf{x}' = H\mathbf{x}$ .

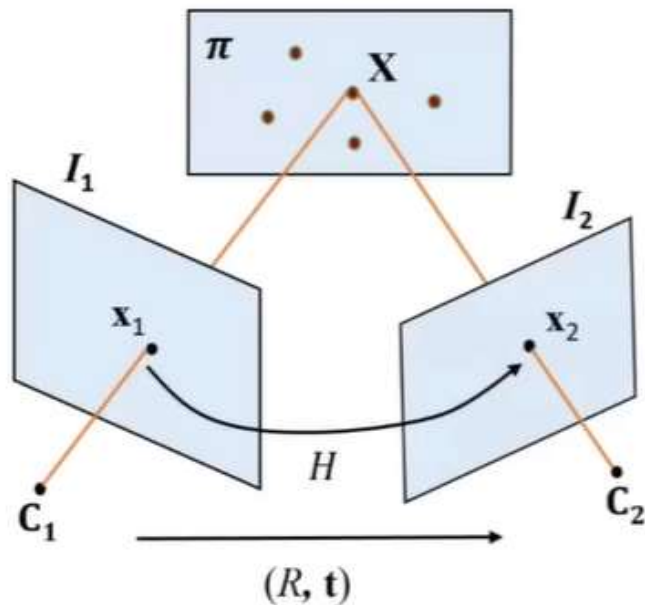


This is also known as  
**Homography!**

Image source: "Multiple View Geometry in Computer Vision", Richard Hartley and Andrew Zisserman

# Existence of Projective Homography

## 1. Planar scene:



- $X_1$  and  $X_2$  is the **3D point  $X$**  expressed in  $C_1$  and  $C_2$  respectively:

$$X_2 = RX_1 + t.$$

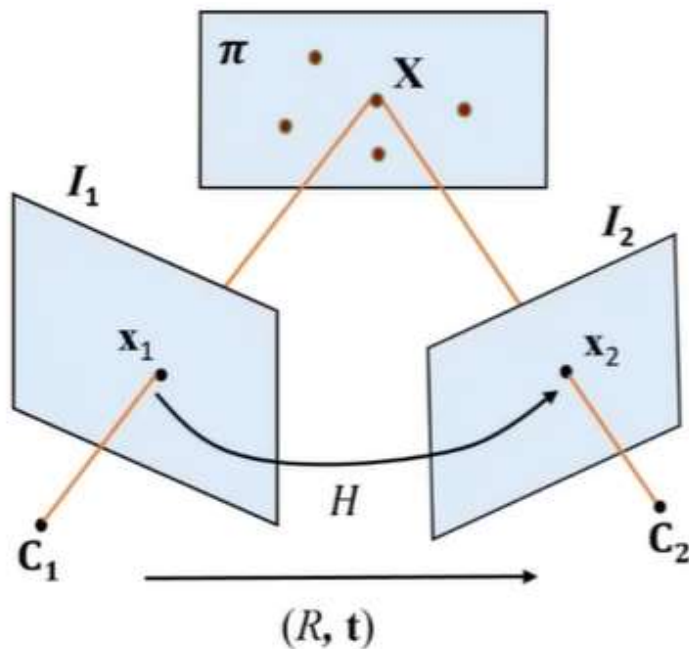
- $N = [n_1, n_2, n_3]^T$  is the **unit normal vector** representing the plane  $\pi$  w.r.t  $C_1$ , and  $d$  is the **perpendicular distance** from plane to  $C_1$ :

$$N^T X_1 = n_1 X + n_2 Y + n_3 Z = d,$$

$$\Rightarrow \frac{N^T X_1}{d} = 1, \quad \forall X_1 \in \pi.$$

# Existence of Projective Homography

## 1. Planar scene:



- Combining the two equations, we get

$$\mathbf{X}_2 = \left( R + \frac{\mathbf{t}\mathbf{N}^T}{d} \right) \mathbf{X}_1,$$

- Since  $\lambda_1 \mathbf{x}_1 = \mathbf{X}_1$  and  $\lambda_2 \mathbf{x}_2 = \mathbf{X}_2$ , we get

$$\lambda \mathbf{x}_2 = \underbrace{\left( R + \frac{\mathbf{t}\mathbf{N}^T}{d} \right)}_H \mathbf{x}_1$$

# Existence of Projective Homography

2. Plane at **infinity**: Scene is very far away from the camera, e.g. aerial images, i.e.

$$H = \left( R + \frac{\mathbf{t}\mathbf{N}^T}{d} \right) \Rightarrow H_{\infty} = \lim_{d \rightarrow \infty} \left( R + \frac{\mathbf{t}\mathbf{N}^T}{d} \right) = R.$$

This is the same as **pure rotation**, i.e.  $\mathbf{t} = (0,0,0)^T$ :

$$H = \left( R + \frac{\mathbf{t}\mathbf{N}^T}{d} \right) \Rightarrow H = R.$$


To continue...

# 2D Homography

- **Given:** A set of **points correspondences**  $\mathbf{x}_i \leftrightarrow \mathbf{x}'_i$  between two images.
- **Compute:** The **2D Homography**,  $H$  such that  $H\mathbf{x}_i = \mathbf{x}'_i$  for each  $i$ .



$\mathbb{P}^2 \rightarrow \mathbb{P}^2$



Point correspondences  
on image planes undergo  
2D Homography



Number of measurements required?

**Question:**

How many corresponding points  $\mathbf{x}_i \leftrightarrow \mathbf{x}'_i$  are required to compute  $H$ ?



# Number of measurements required?

## Answer:

- The number of **degrees of freedom** and number of **constraints** give a lower bound:
  1. **8 degrees of freedom** for  $H$ , i.e. 9 entries less 1 for up to scale.
  2. We will see that each point correspondence  $\mathbf{x}_i \leftrightarrow \mathbf{x}'_i$  gives **2 constraints**.
- As a consequence, it is necessary to specify **four point correspondences** in order to constrain  $H$  fully.

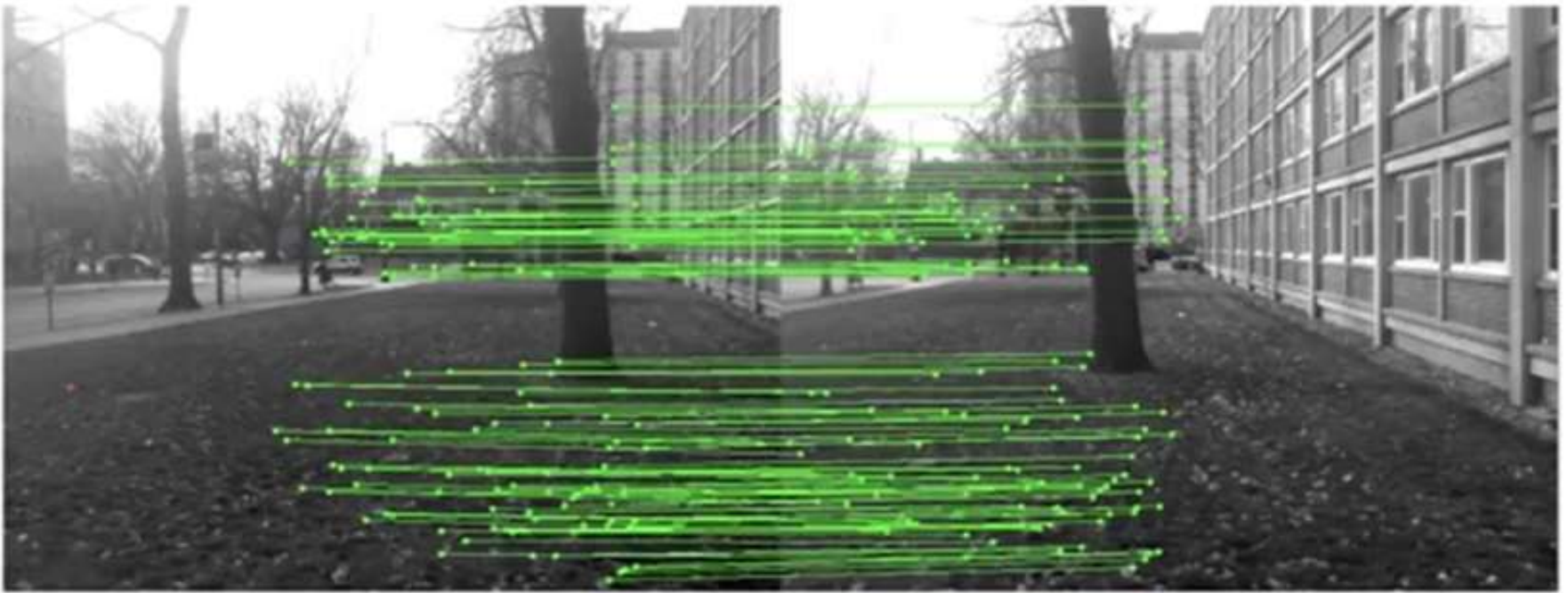
# Approximate Solution

- It will be seen that if exactly **four correspondences** are given, then **an exact solution** for the matrix  $H$  is possible.
- This is the **minimal solution**, which is important for the **number of RANSAC loops** for robust estimation (details later).
- Since points are measured inexactly (“noise”), more than four correspondences are usually used to obtain a **least-squares solution** (details later).

# Possible ways

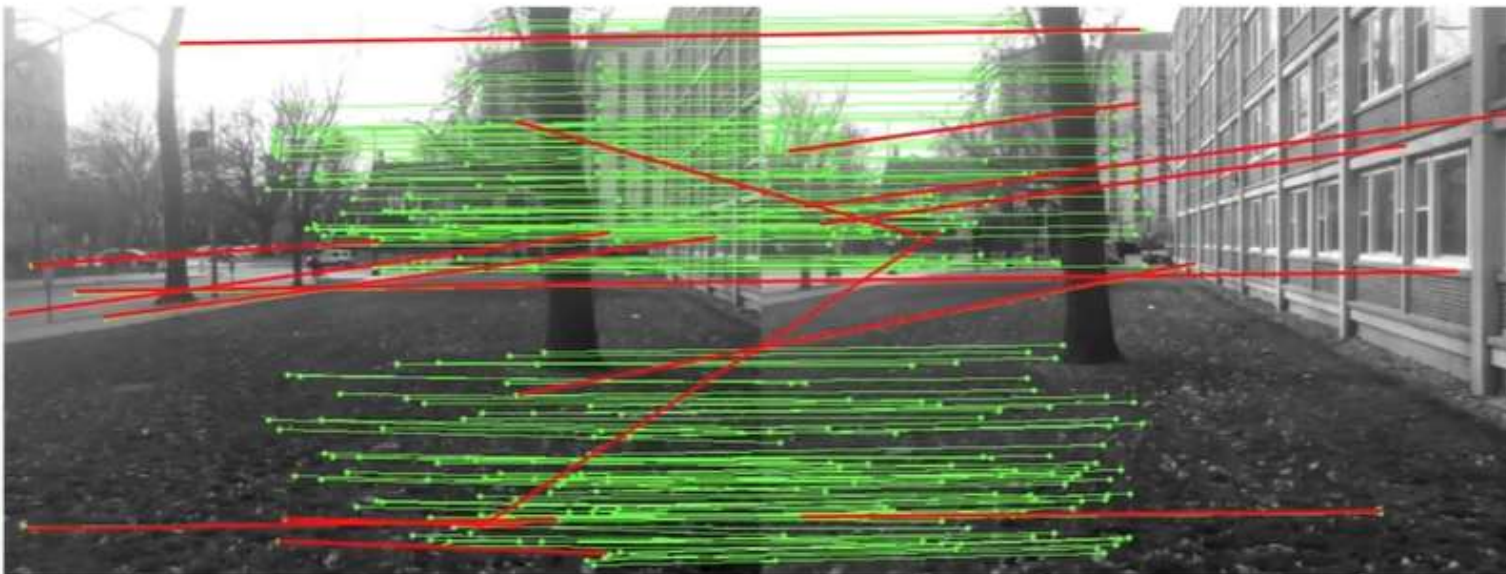
- Direct Linear Transformation (DLT) Algorithm
  - Least square method
  - Iterative method
- RANSAC Algorithm

# RANdom SAmple Consensus: RANSAC



# RANdom SAmple Consensus: RANSAC

- In reality, keypoint matching gives us many **outliers**.
- Outliers can **severely disturb** the least-squares estimation, and should be removed.





# The RANSAC Song

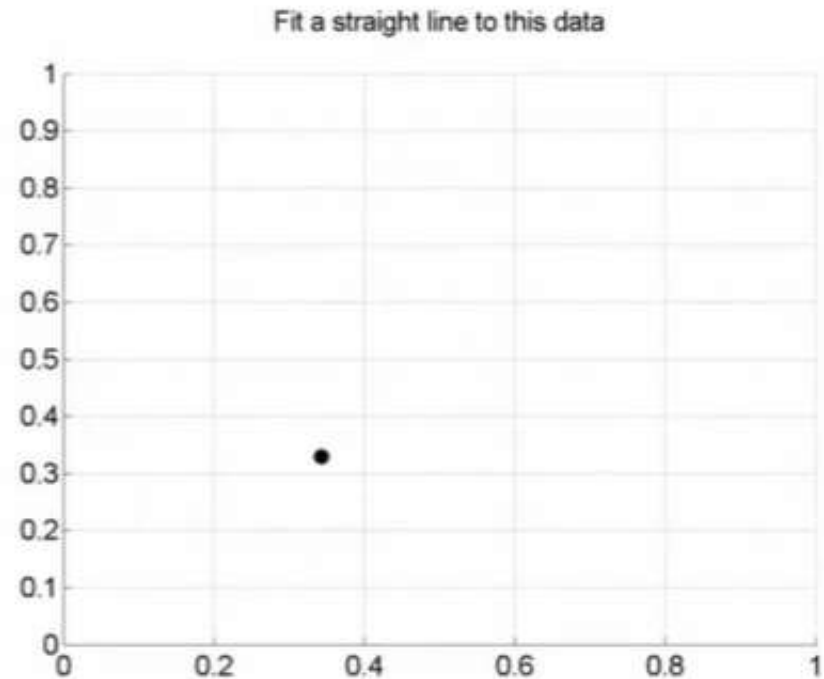
When you have outliers you may face much frustration  
if you include them in a model fitting operation.  
But if your model's fit to a sample set of minimal size,  
the probability of the set being outlier-free will rise.  
Brute force tests of all sets will cause computational constipation.

$N$  random samples  
will provide an example  
of a fitted model uninfluenced by outliers. No need to test all combinations!

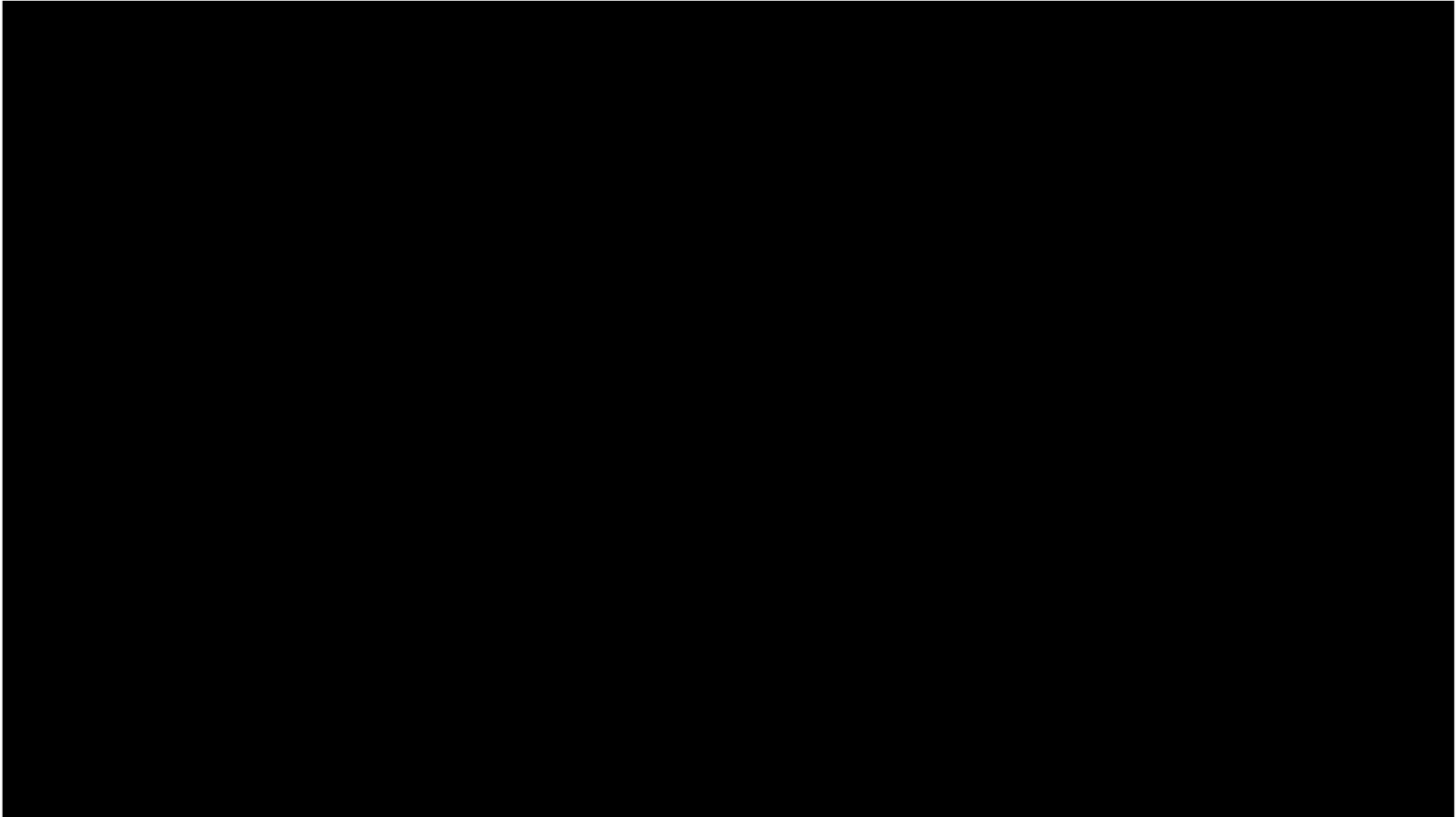
Each random trial should have its own unique sample set  
and make sure that the sets you choose are not degenerate.  
 $N$ , the number of sets, to choose is based on the probability  
of a point being an outlier, and of finding a set that's outlier free.  
Updating  $N$  as you go will minimise the time spent.

So if you gamble  
that  $N$  samples are ample  
to fit a model to your set of points, it's likely that you will win the bet.

Select the set that boasts  
that its number of inliers is the most (you're almost there).  
Fit a new model just to those inliers and discard the rest,  
an estimated model for your data is now possessed!  
This marks the end point of your model fitting quest.

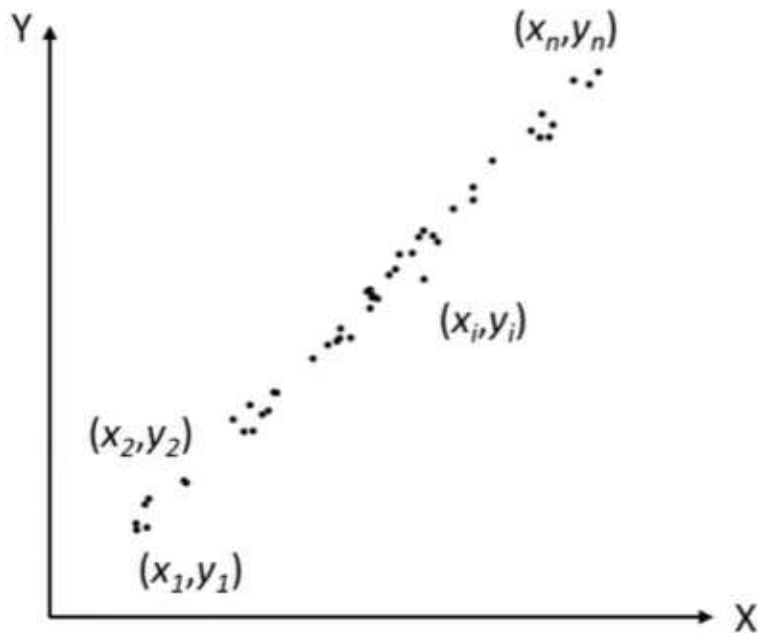


# The RANSAC Song



# RANSAC: Line fitting example

- **Given:**  $n$  data points  $(x_i, y_i)$ , for  $i = 1, \dots, n$
- **Find:** Best fit line, i.e. **two parameters**  $(m, c)$  from the line equation  $y_i = mx_i + c$ , for  $i = 1, \dots, n$

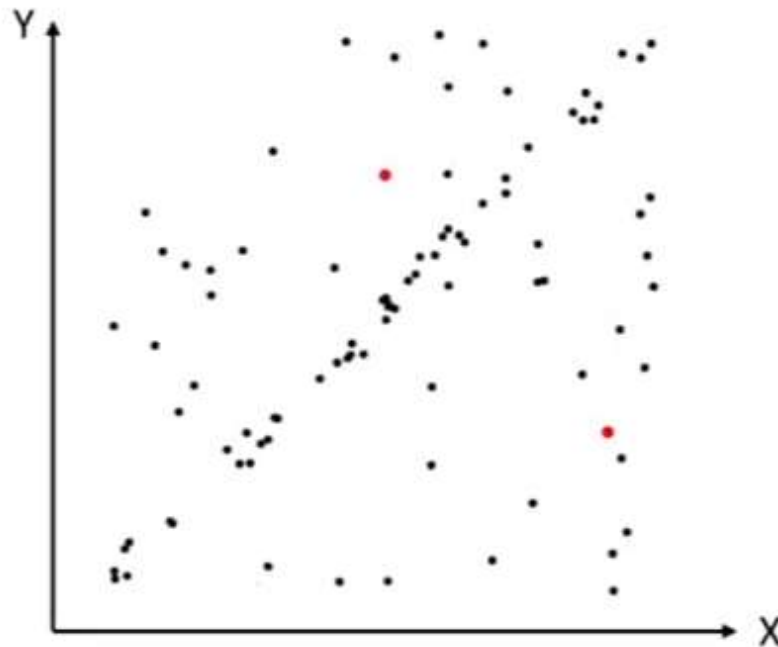




# RANSAC: Line fitting example

## RANSAC Steps:

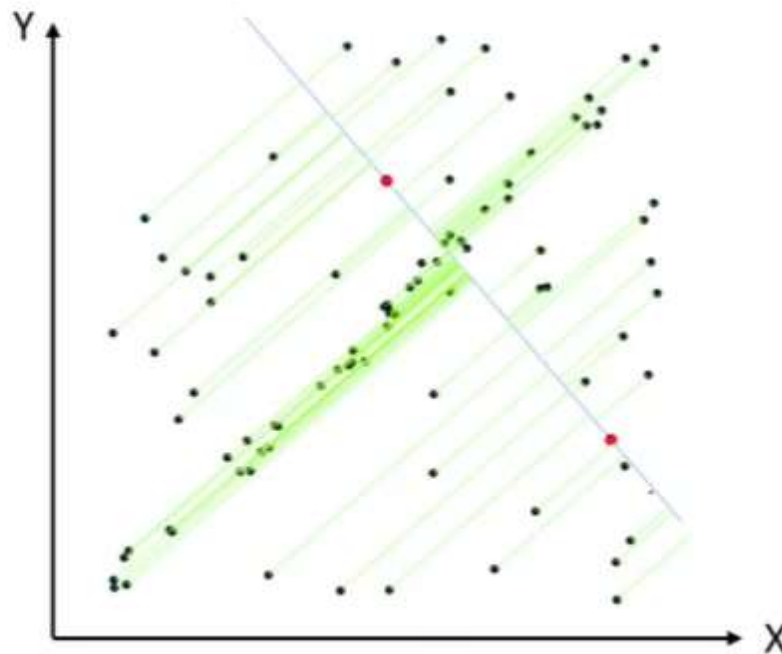
1. Randomly select **minimal subset** of points, i.e. 2 points



# RANSAC: Line fitting example

## RANSAC Steps:

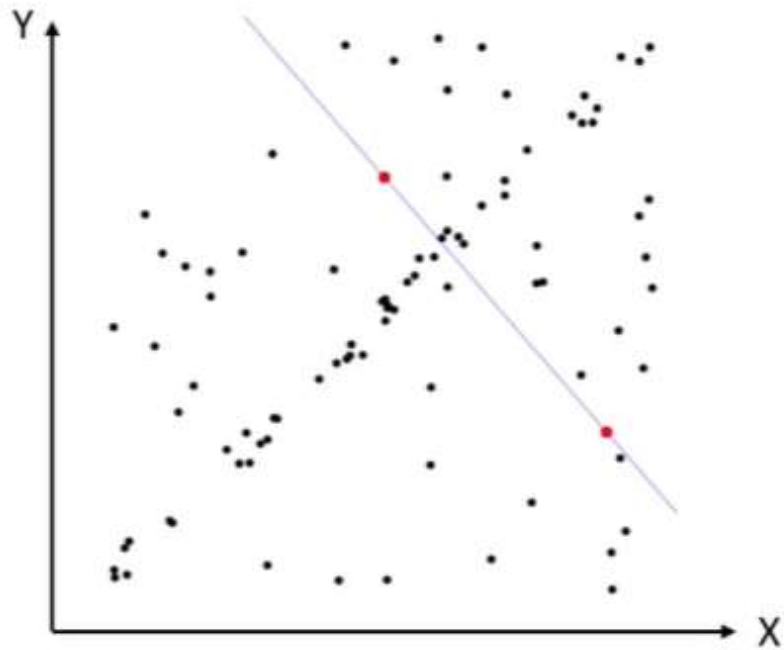
3. Compute **error function**, i.e. shortest point to line distance



# RANSAC: Line fitting example

## RANSAC Steps:

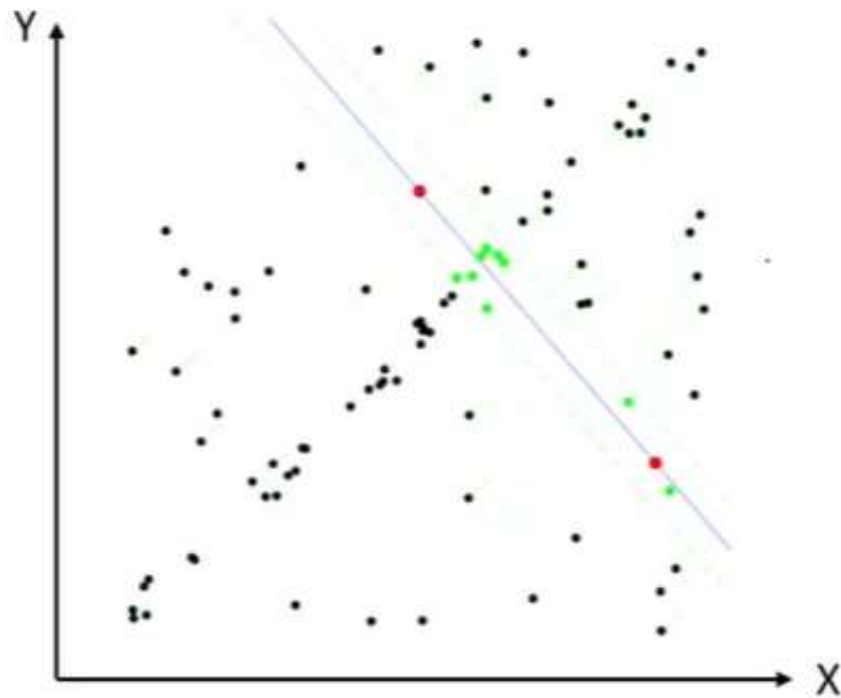
2. Hypothesize a model



# RANSAC: Line fitting example

## RANSAC Steps:

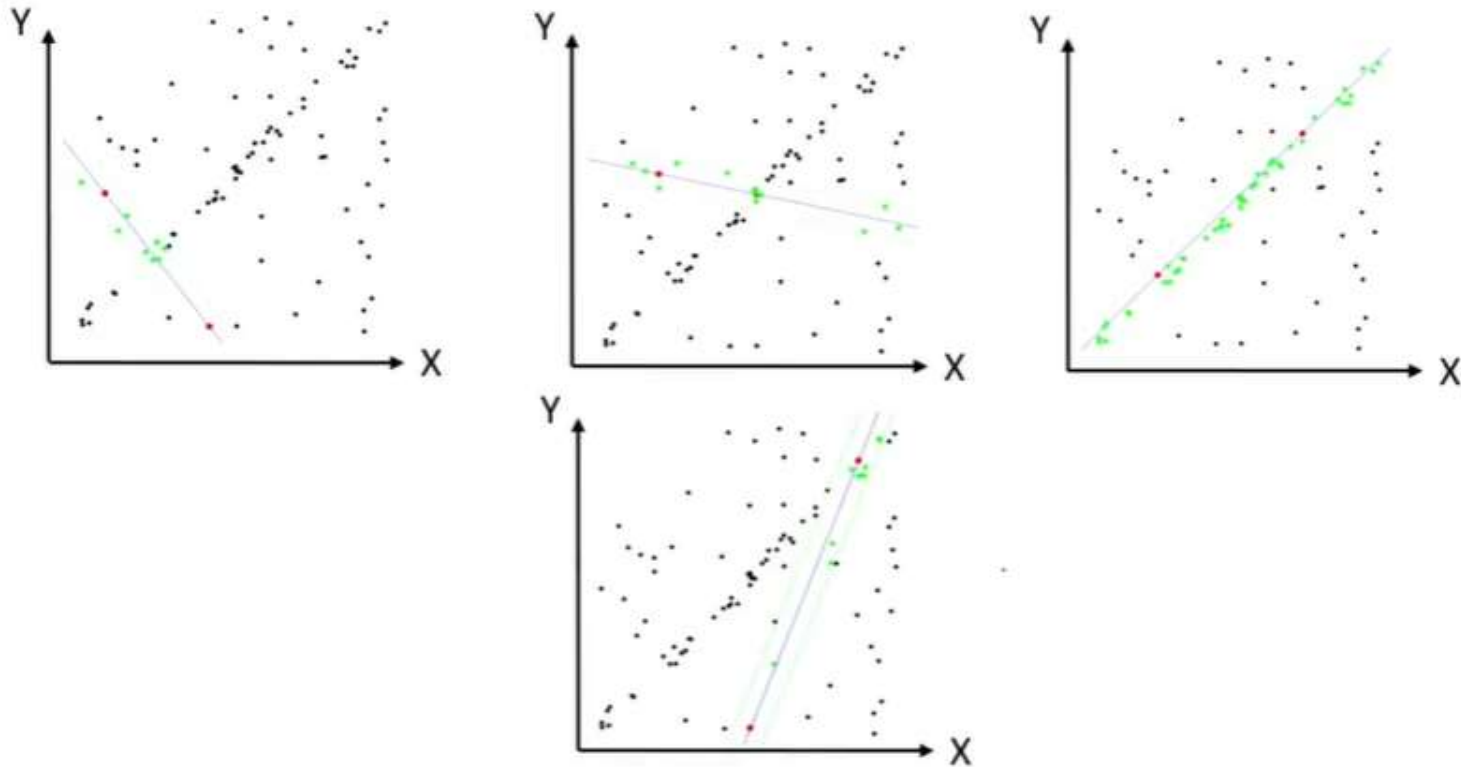
4. Select points **consistent** with model



# RANSAC: Line fitting example

## RANSAC Steps:

5. **Repeat** hypothesize-and-verify loop



# RANSAC Algorithm

## Objective

Robust fit of a model to a data set  $S$  which contains outliers.

## Algorithm

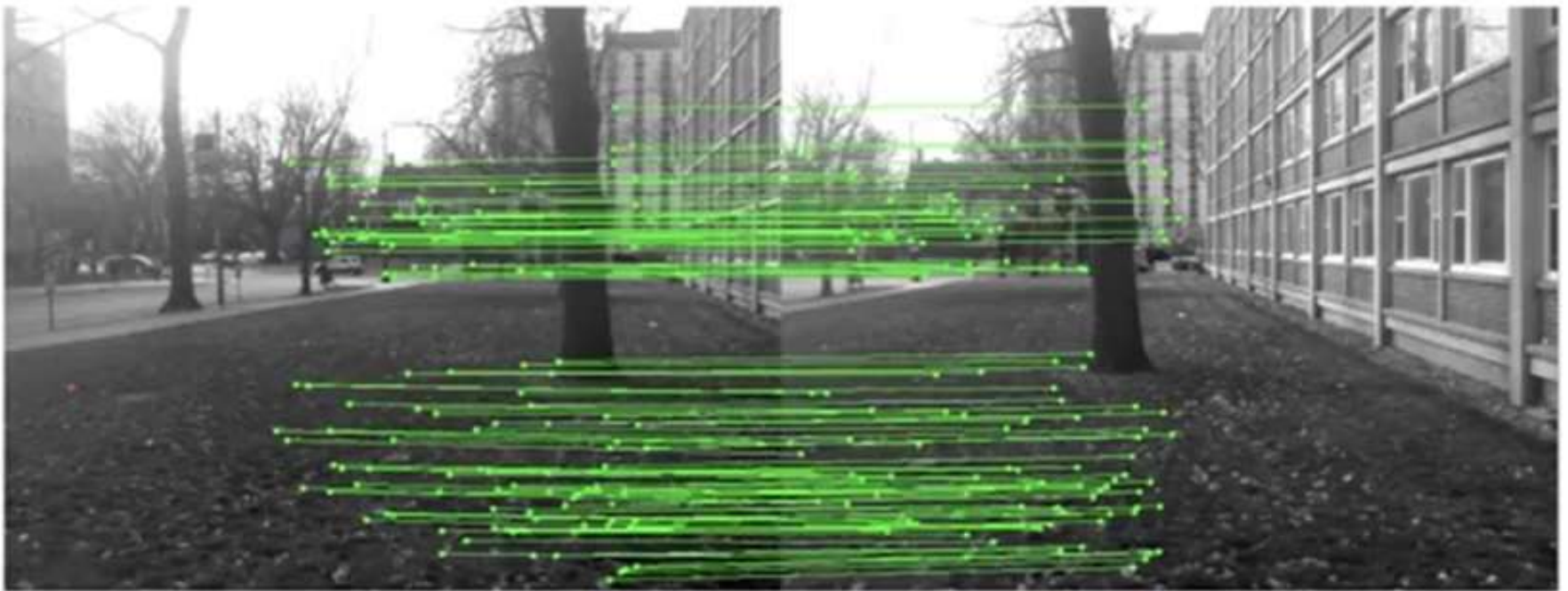
- i. **Randomly select** a sample of  $s$  data points from  $S$  and instantiate the model from this subset.
- ii. Determine the set of data points  $S_i$  which are **within a distance threshold  $t$**  of the model. The set  $S_i$  is the **consensus set** of the sample and defines the inliers of  $S$ .
- iii. After  $N$  trials, select the **largest consensus set  $S_i$** . The model is re-estimated using all the points in the subset  $S_i$ .

Three parameters:

Number of points	$s$
Distance threshold	$t$
Number of Samples	$N$

M. Fischler, R. Bolles, "Random sample consensus: a paradigm for model fitting with applications to image analysis and automated cartography", Communications ACM, 1981.

How to find the points of interest?



To continue...