# Lecture 15 Robust Estimation: RANSAC

Let's say we have found point matches between two images, and we think they are related by some parametric transformation (e.g. translation; scaled Euclidean; affine). How do we estimate the parameters of that transformation?

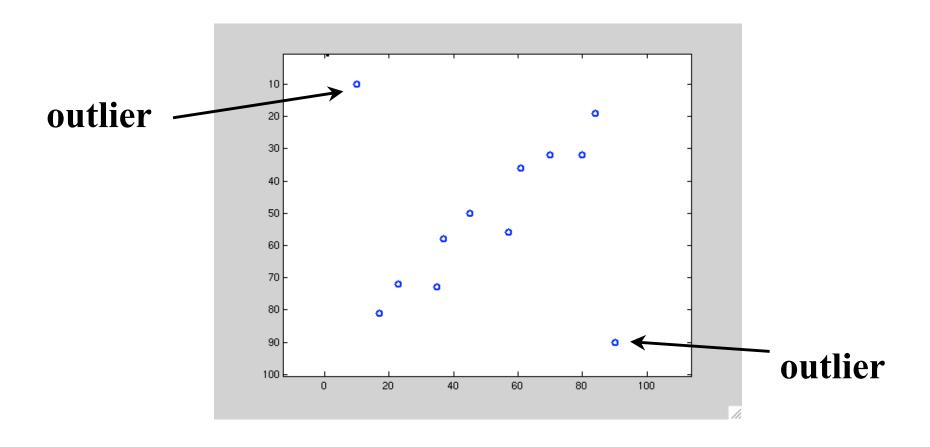
#### **General Strategy**

• Least-Squares estimation from point correspondences

But there are problems with that approach....

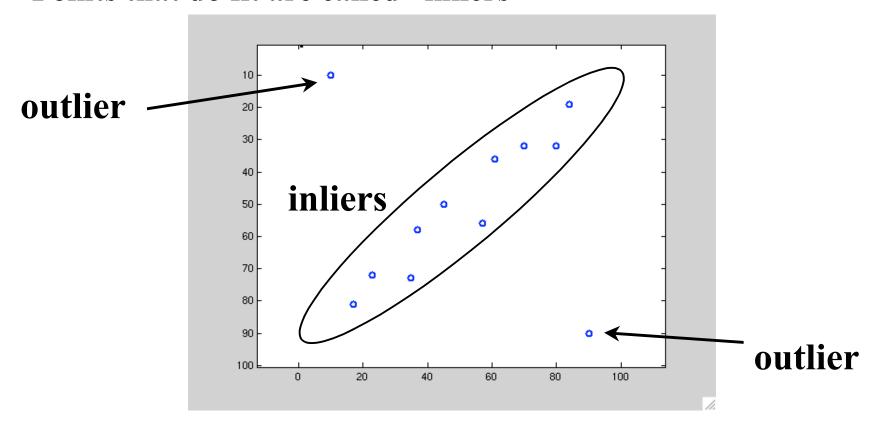
#### **Problem: Outliers**

Loosely speaking, outliers are points that don't "fit" the model.



#### **Bad Data => Outliers**

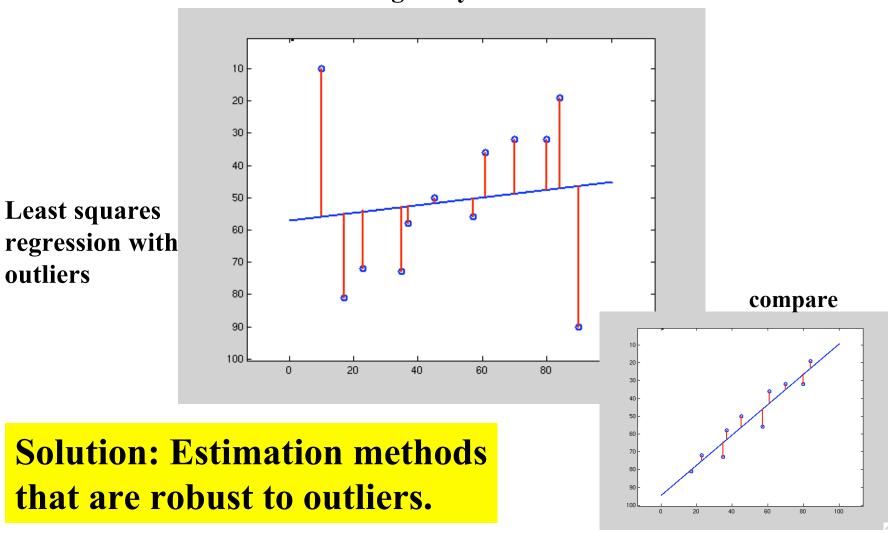
Loosely speaking, outliers are points that don't "fit" the model. Points that do fit are called "inliers"

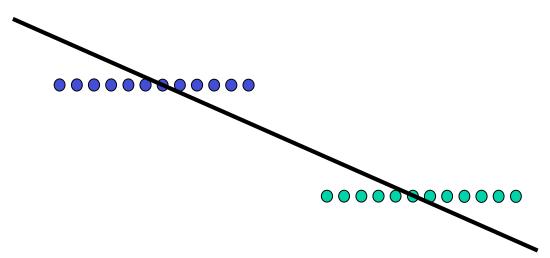


outliers

#### **Problem with Outliers**

Least squares estimation is sensitive to outliers, so that a few outliers can greatly skew the result.



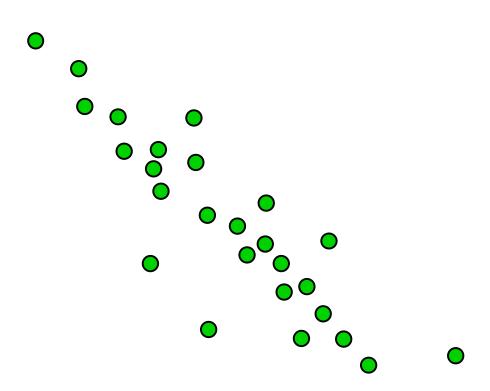


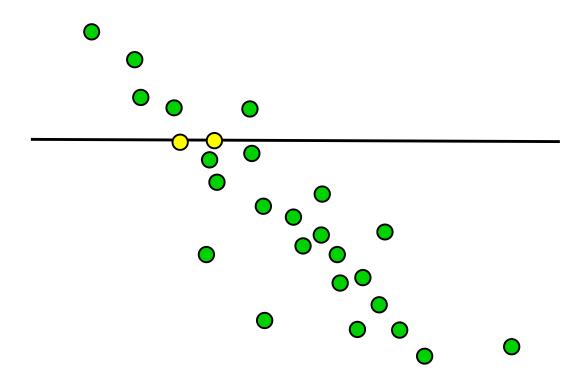
Multiple structures can also skew the results. (the fit procedure implicitly assumes there is only one instance of the model in the data).

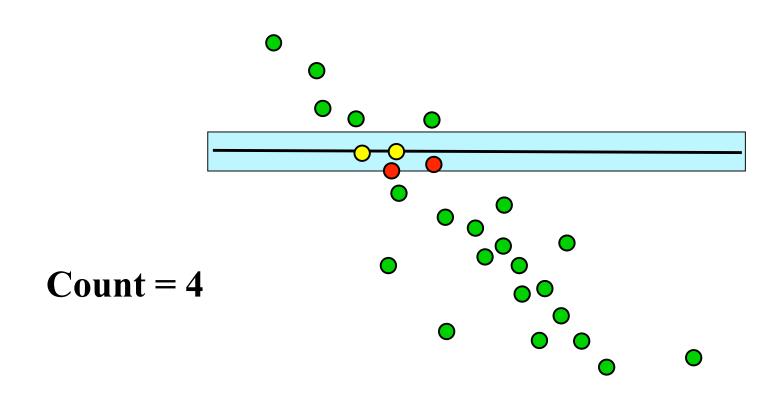
#### **Robust Estimation**

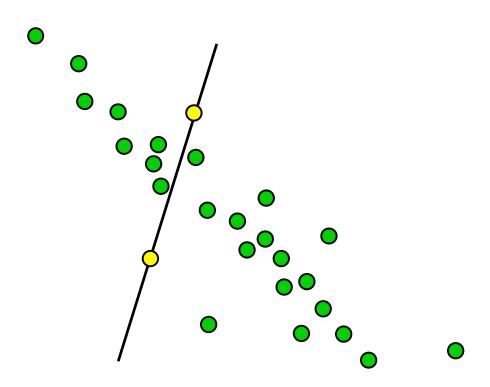
- View estimation as a two-stage process:
  - Classify data points as outliers or inliers
  - Fit model to inliers while ignoring outliers
- Example technique: RANSAC
   (RANdom SAmple Consensus)

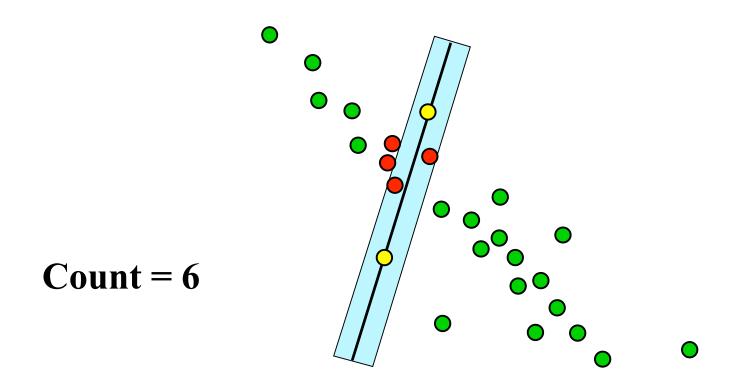
M. A. Fischler and R. C. Bolles (June 1981). "Random Sample Consensus: A Paradigm for Model Fitting with Applications to Image Analysis and Automated Cartography". *Comm. of the ACM* **24**: 381--395.

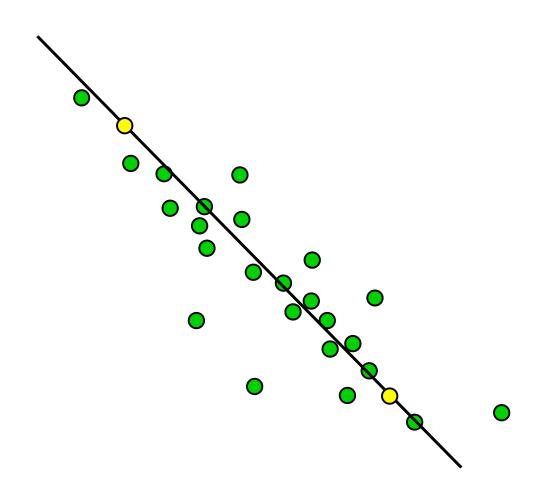


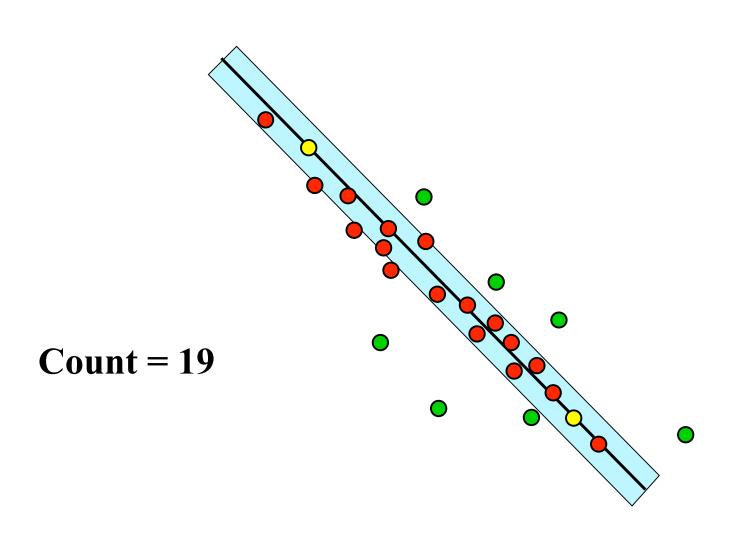


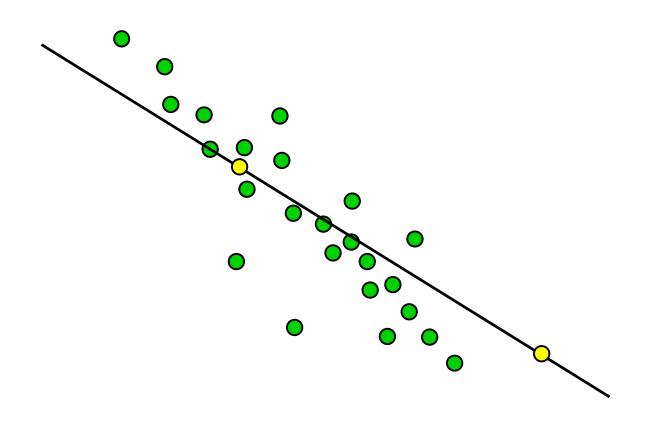


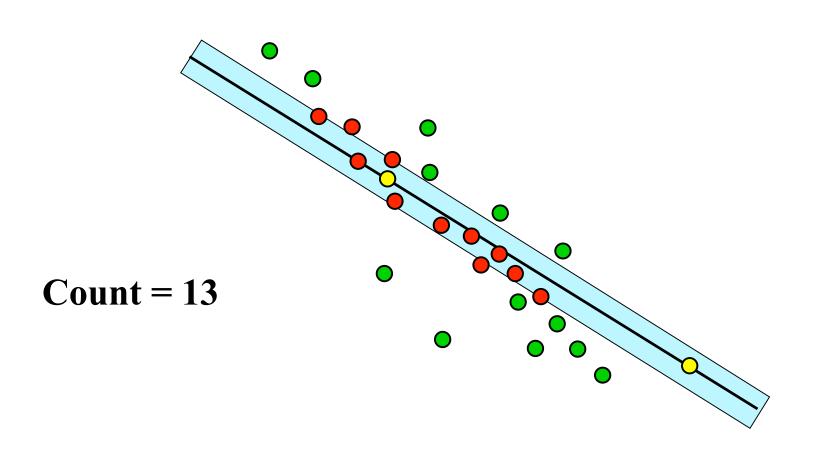


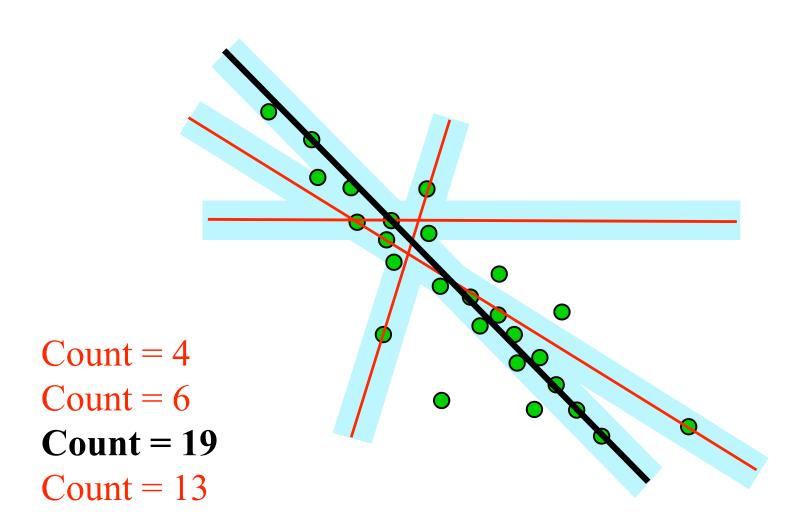












#### Algorithm 15.4: RANSAC: fitting lines using random sample consensus

```
Determine:
    S — the smallest number of points required
    N— the number of iterations required
    d— the threshold used to identify a point that fits well
    T— the number of nearby points required
      to assert a model fits well
Until Niterations have occurred
    Draw a sample of S points from the data
      uniformly and at random
    Fit to that set of S points
    For each data point outside the sample
       Test the distance from the point to the line
         against d if the distance from the point to the line
         is less than d the point is close
    end
    If there are T or more points close to the line
      then there is a good fit. Refit the line using all
      these points.
end
Use the best fit from this collection, using the
  fitting error as a criterion
```

#### (Forsyth & Ponce)

e = probability that a point is an outlier

s = number of points in a sample

N = number of samples (we want to compute this)

p = desired probability that we get a good sample

#### Solve the following for N:

$$1 - (1 - (1 - e)^{s})^{N} = p$$

Where in the world did that come from? ....

e = probability that a point is an outlier

s = number of points in a sample

N = number of samples (we want to compute this)

p = desired probability that we get a good sample

$$1 - (1 - (1 - e)^{s})^{N} = p$$

**Probability that choosing** one point yields an inlier

e = probability that a point is an outlier

s = number of points in a sample

N = number of samples (we want to compute this)

p = desired probability that we get a good sample

$$1 - (1 - (1 - e)^{s})^{N} = p$$

Probability of choosing s inliers in a row (sample only contains inliers)

e = probability that a point is an outlier

s = number of points in a sample

N = number of samples (we want to compute this)

p = desired probability that we get a good sample

$$1 - (1 - (1 - e)^{s})^{N} = p$$

Probability that one or more points in the sample were outliers (sample is contaminated).

e = probability that a point is an outlier

s = number of points in a sample

N = number of samples (we want to compute this)

p = desired probability that we get a good sample

$$1 - (1 - (1 - e)^{s})^{N} = p$$

**Probability that N samples** were contaminated.

e = probability that a point is an outlier

s = number of points in a sample

N = number of samples (we want to compute this)

p = desired probability that we get a good sample

$$1 - (1 - (1 - e)^{s})^{N} = p$$

Probability that at least one sample was not contaminated (at least one sample of s points is composed of only inliers).

## How many samples?

Choose N so that, with probability p, at least one random sample is free from outliers. e.g. p=0.99

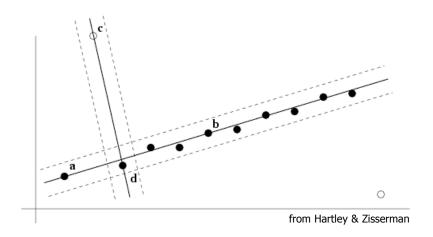
$$(1 - (1 - e)^s)^N = 1 - p$$

$$N = \frac{\log(1-p)}{\log(1-(1-e)^s)}$$

	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		

# CSE486, Penn State Example: N for the line-fitting problem

- n = 12 points
- Minimal sample size S = 2
- 2 outliers:  $e = 1/6 \Rightarrow 20\%$
- So N = 5 gives us a 99% chance of getting a pure-inlier sample
  - Compared to N = 66 by trying every pair of points



## Acceptable consensus set?

- We have seen that we don't have to exhaustively sample subsets of points, we just need to randomly sample N subsets.
- However, typically, we don't even have to sample N sets!
- <u>Early termination</u>: terminate when inlier ratio reaches expected ratio of inliers

$$T = (1 - e) * (total number of data points)$$

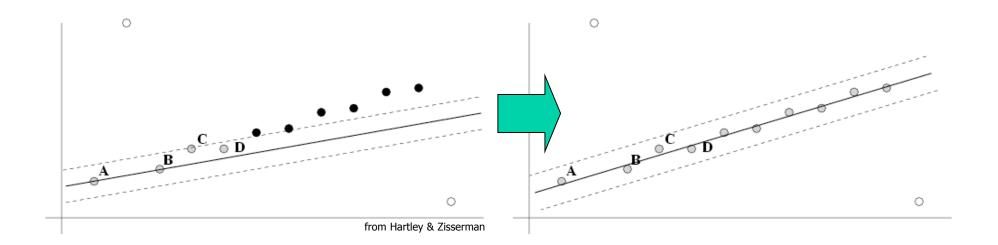
## RANSAC: Picking Distance Threshold C

- Usually chosen empirically
- But...when measurement error is known to be Gaussian with mean 0 and variance  $S^2$ :
  - Sum of squared errors follows a  $\chi^2$  distribution with  $\mathbf{m}$  DOF, where  $\mathbf{m}$  is the DOF of the error measure (the *codimension*)
  - (dimension + codimension) = dimension of parameter space
    - E.g., m = 1 for line fitting because error is perpendicular distance
    - E.g., m = 2 for point distance
- Examples for probability p = 0.95 that point is inlier

m	Model	$d^2$	
1	Line, fundamental matrix	$3.84  \mathrm{S}^2$	
2	Homography, camera matrix	5.99 <b>S</b> <sup>2</sup>	

#### **After RANSAC**

- RANSAC divides data into inliers and outliers and yields estimate computed from minimal set of inliers with greatest support
- Improve this initial estimate with Least Squares estimation over all inliers (i.e., standard minimization)
- Find inliers wrt that L.S. line, and compute L.S. one more time.



## **Practical Example**

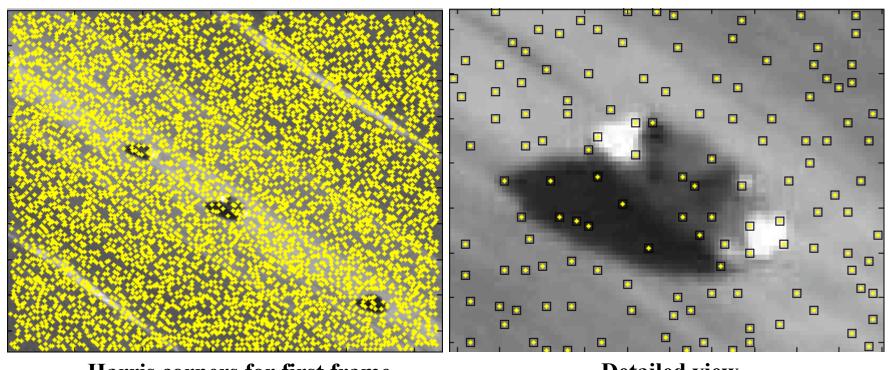
- Stabilizing aerial imagery using RANSAC
  - find corners in two images
  - hypothesize matches using NCC
  - do RANSAC to find matches consistent with an affine transformation
  - take the inlier set found and estimate a full projective transformation (homography)

Input: two images from an aerial video sequence.



Note that the motion of the camera is "disturbing"

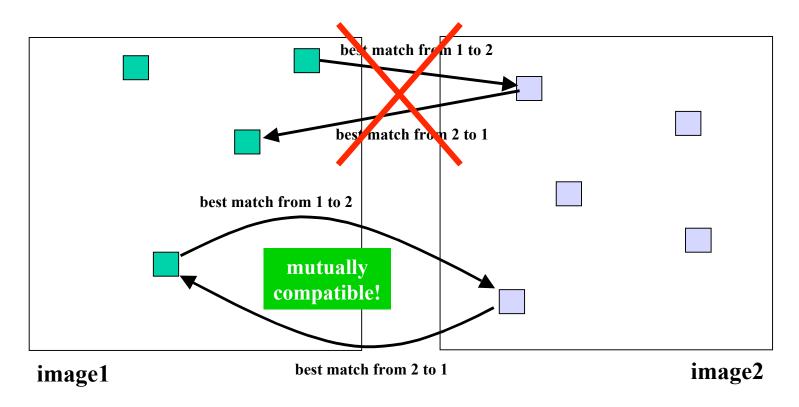
Step1: extract Harris corners from both frames. We use a small threshold for R because we want LOTS of corners (fodder for our next step, which is matching).



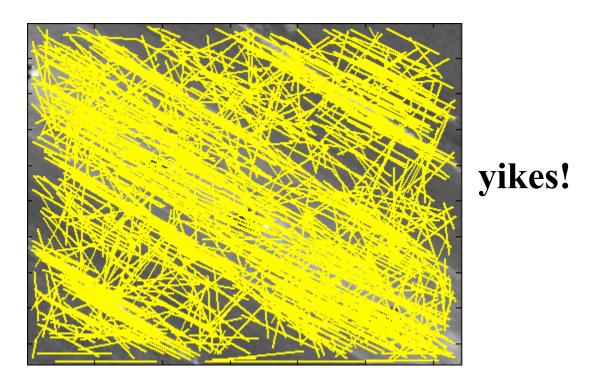
Harris corners for first frame

**Detailed view** 

Step2: hypothesize matches. For each corner in image 1, look for matching intensity patch in image2 using NCC. Make sure matching pairs have highest NCC match scores in BOTH directions.

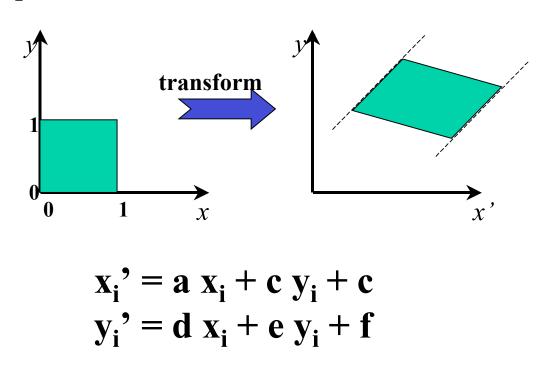


Step2: hypothesize matches.



As you can see, a lot of false matches get hypothesized. The job of RANSAC will be to clean this mess up.

Step3: Use RANSAC to robustly fit best affine transformation to the set of point matches.



How many unknowns? How many point matches are needed?

Step3: Use RANSAC to robustly fit best affine transformation to the set of point matches.

Affine transformation has 6 degrees of freedom. We therefore need 3 point matches [each gives 2 equations]

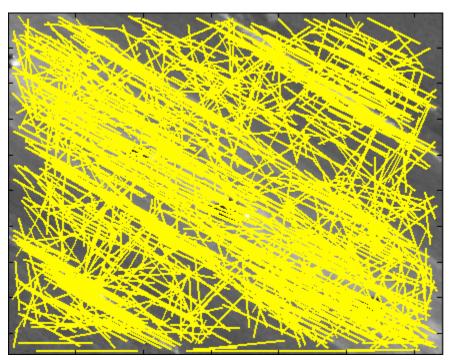
Randomly sample sets of 3 point matches. For each, compute the unique affine transformation they define. How?

How to compute affine transformation from 3 point matches? Use Least Squares! (renewed life for a nonrobust approach)

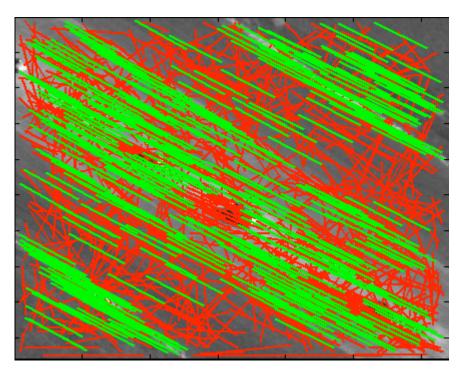
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Then transform all points from image1 to image2 using that computed transformation, and see how many other matches confirm the hypothesis.

Repeat N times.



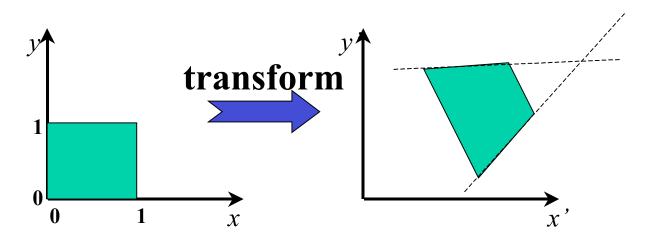
original point matches



labels from RANSAC green: inliers

red: outliers

Step4: Take inlier set labeled by RANSAC, and now use least squares to estimate a projective transformation that aligns the images. (we will discuss this ad nauseum in a later lecture).



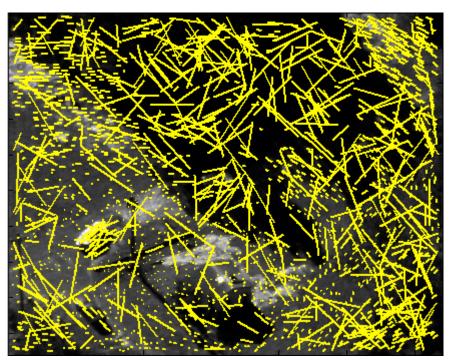
**Projective Transformation** 

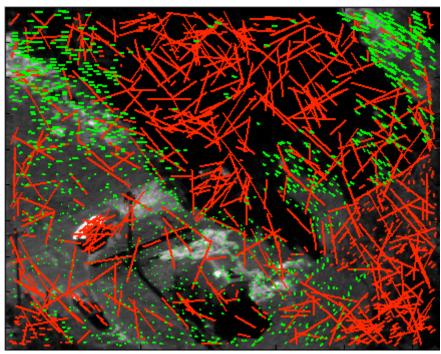
Step4: estimate projective transformation that aligns the images.



Now it is easier for people (and computers) to see the moving objects.





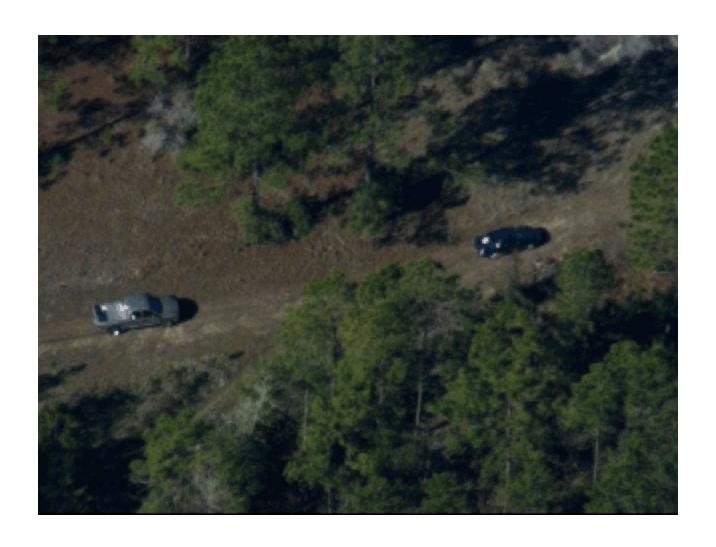


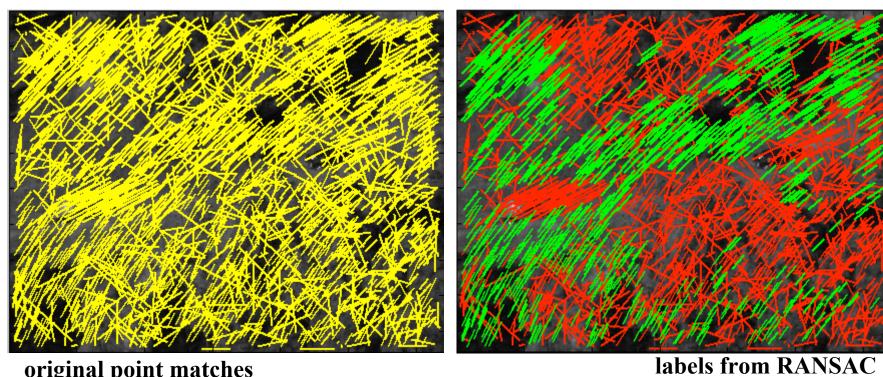
original point matches

labels from RANSAC

green: inliers red: outliers







original point matches

green: inliers red: outliers

