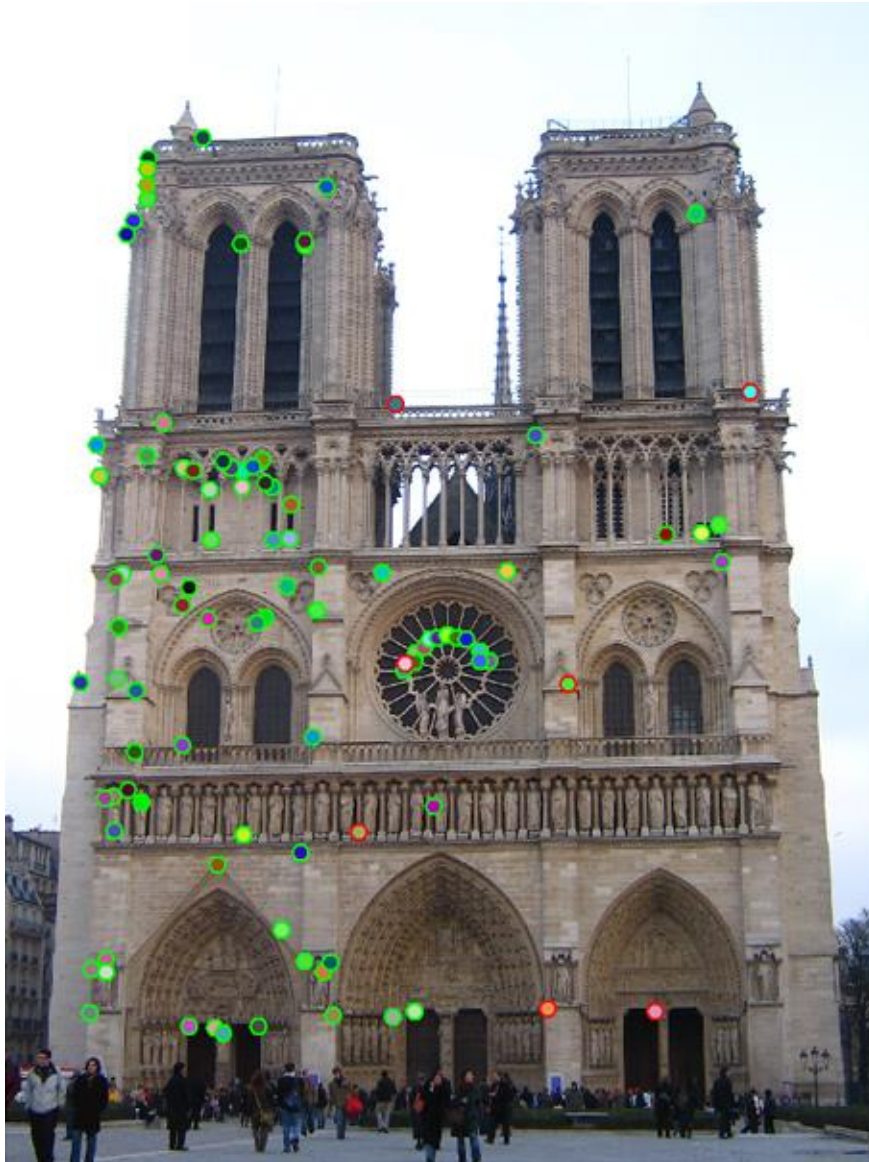


An abstract green geometric pattern consisting of a network of interconnected points and lines, resembling a mesh or a complex polygonal structure, set against a dark green background.

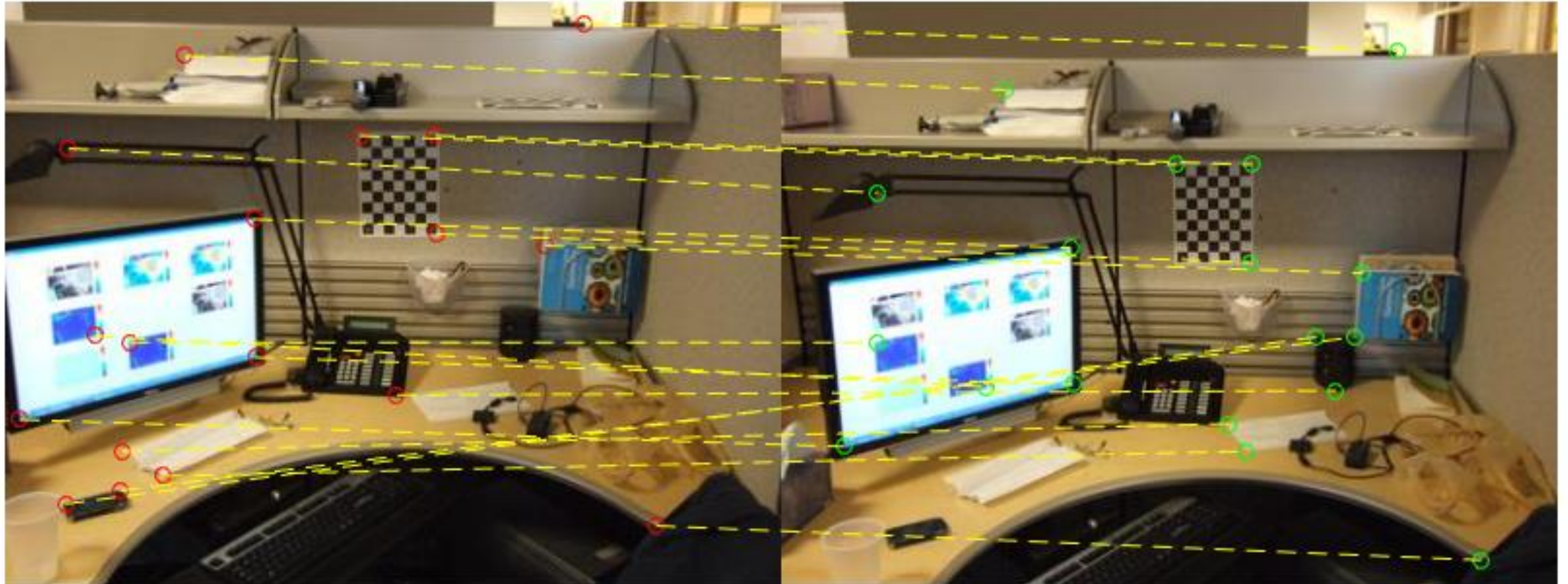
RANSAC

Matching Points

Given matches, what Next?



Stereo images with outliers



Stereo images without outliers

Point matches after outliers were removed



Fitting and Alignment

Fitting :

Find the parameters of a model that best fit the data.

A l i g n m e n t :

Find the parameters of the transformation that best aligns matched points.

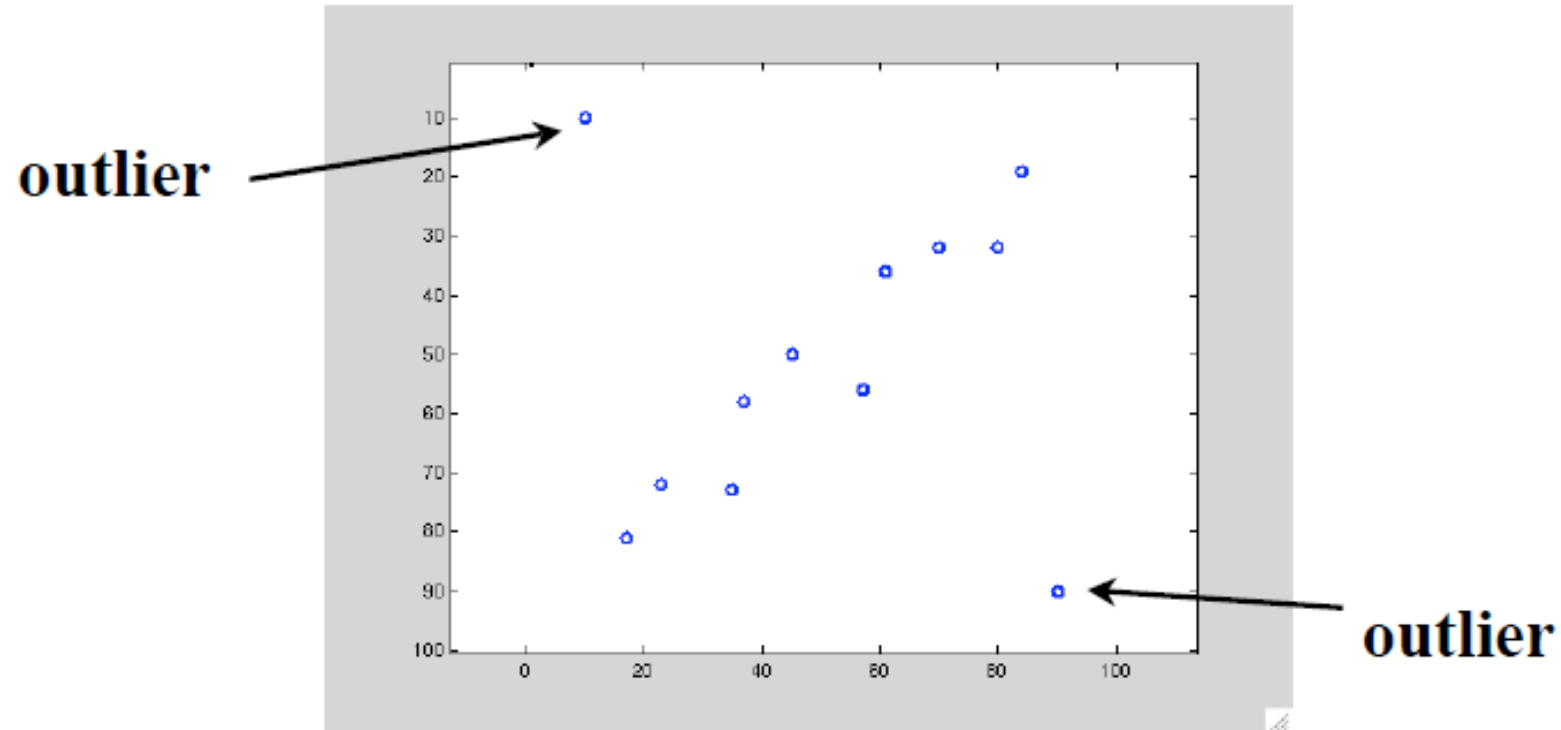
G e n e r a l S t r a t e g y :

- Least-Squares estimation from point correspondences

But there are problems with that approach....

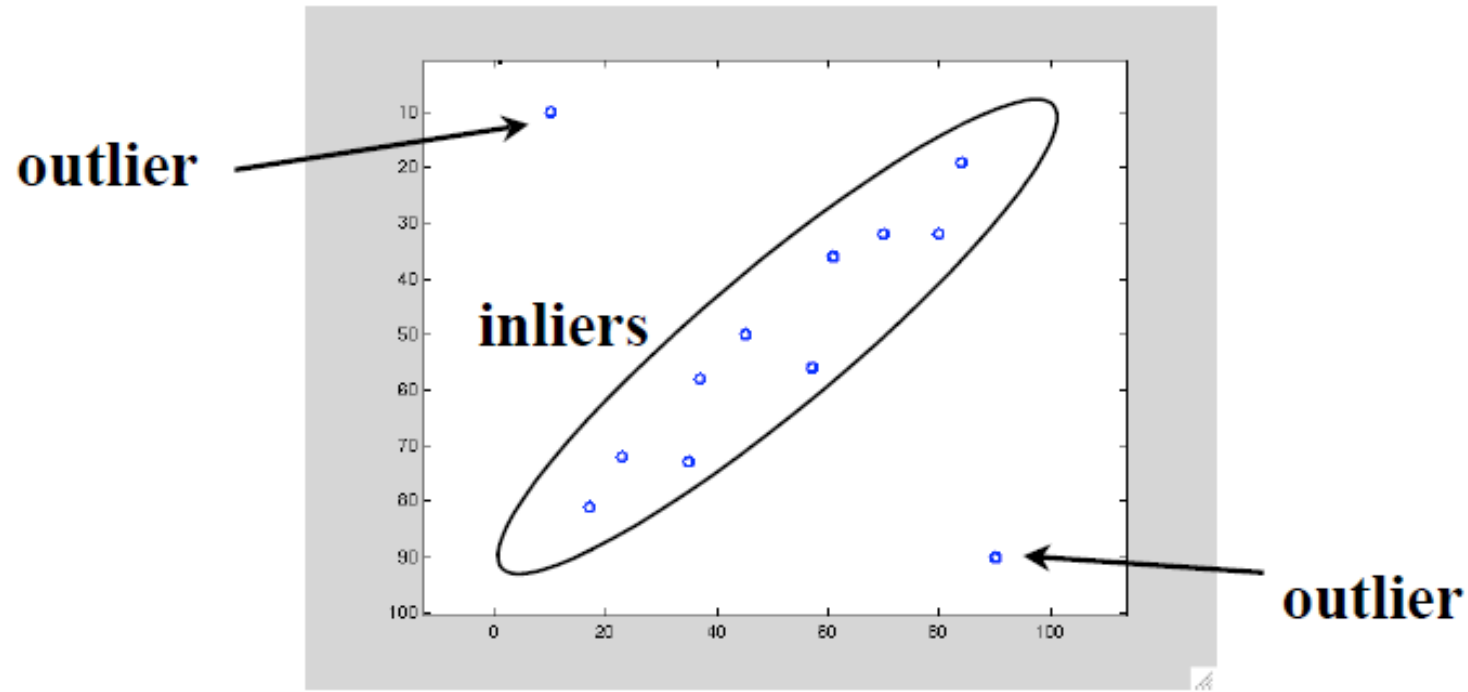
Problem : Outliers

- Loosely speaking, outliers are points that don't “fit” the model.



Bad Data => Outliers

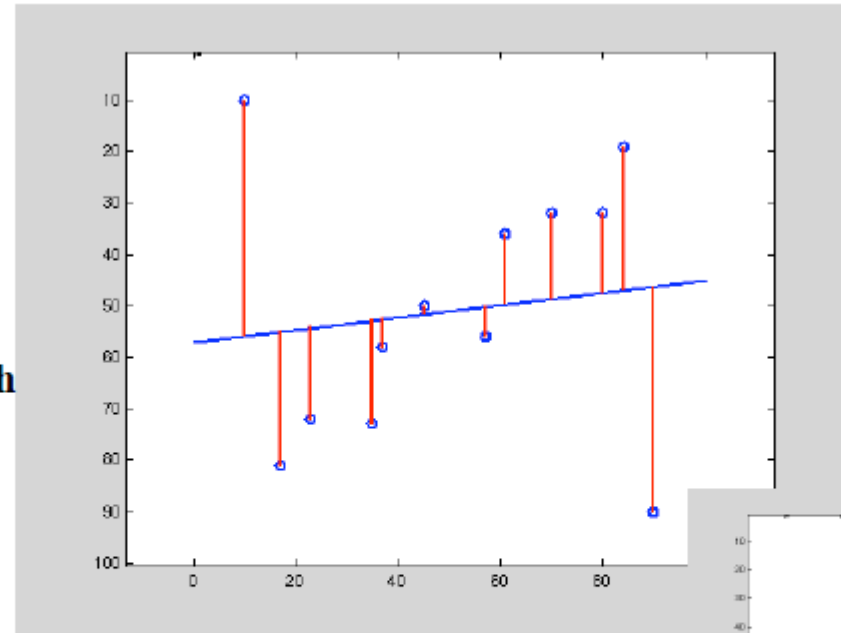
- Loosely speaking, outliers are points that don't "fit" the model. Points that do fit are called "inliers"



Problem with Outliers

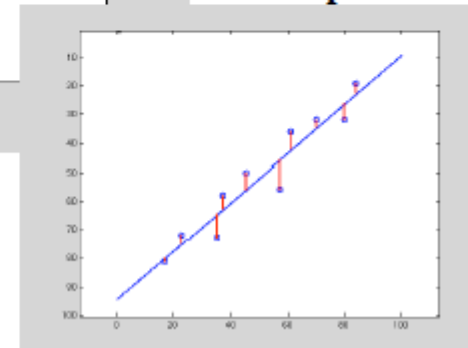
- Least squares estimation is sensitive to outliers, so that a few outliers can greatly skew the result.

Least squares
regression with
outliers



**Solution: Estimation methods
that are robust to outliers.**

compare



Robust Estimation

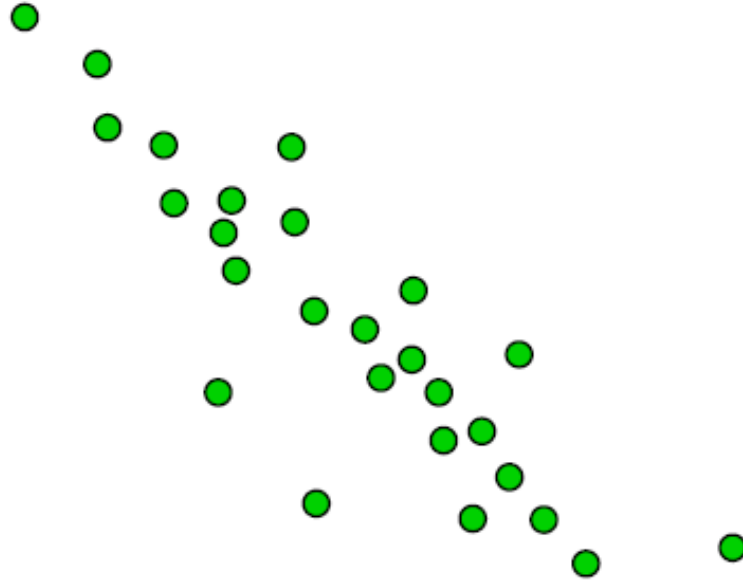
- View estimation as a two-stage process:
 - Classify data points as outliers or inliers
 - Fit model to inliers while ignoring outliers

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.

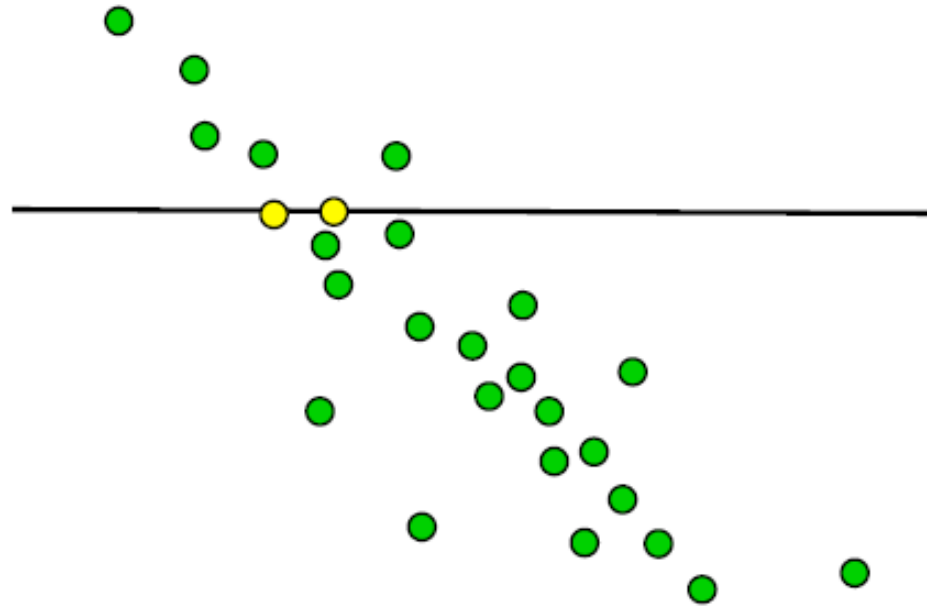
RANSAC

- **RAN**dom **SA**mples **C**onsensus
- Approach: we want to avoid the impact of outliers, so let's look for “inliers”, and use only those.
- Intuition: if an outlier is chosen to compute the current fit, then the resulting line won't have much support from rest of the points.

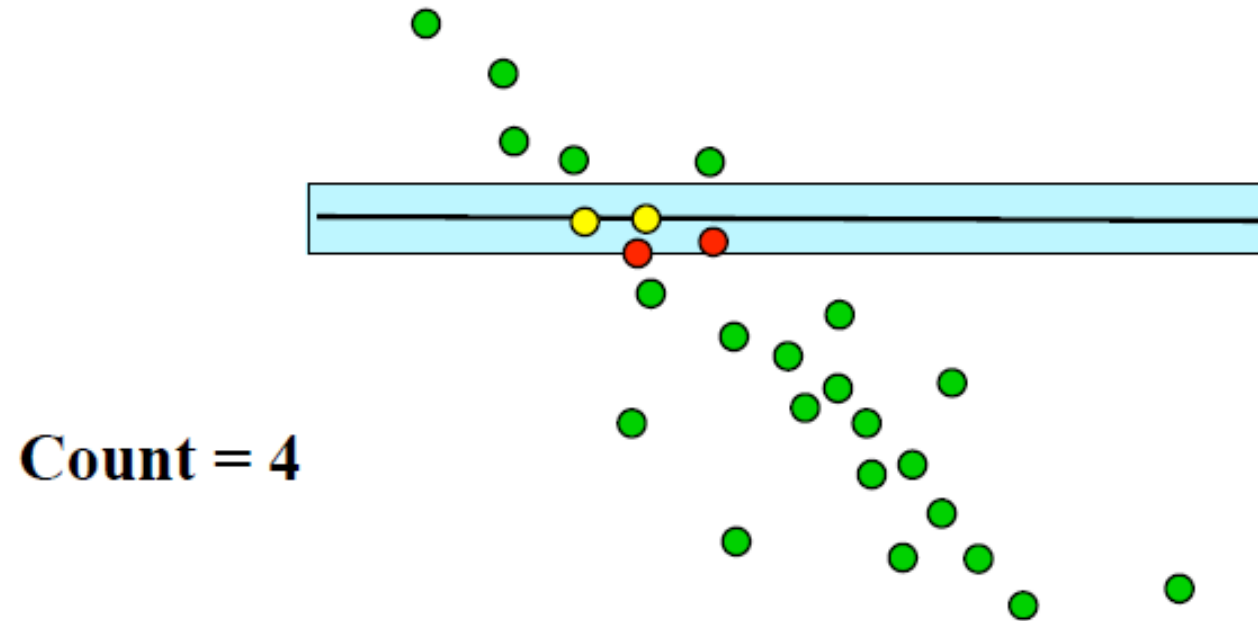
RANSAC Procedure



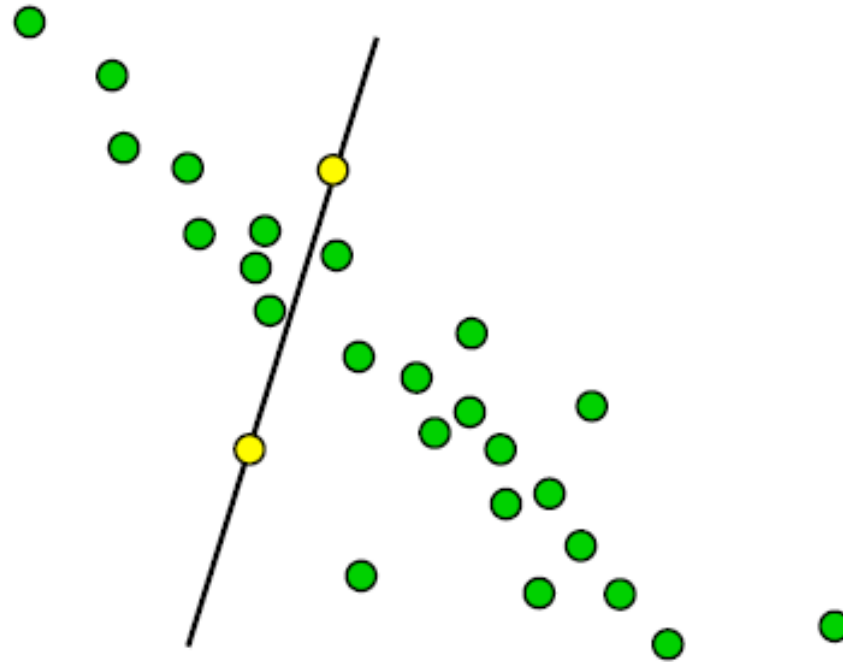
RANSAC Procedure



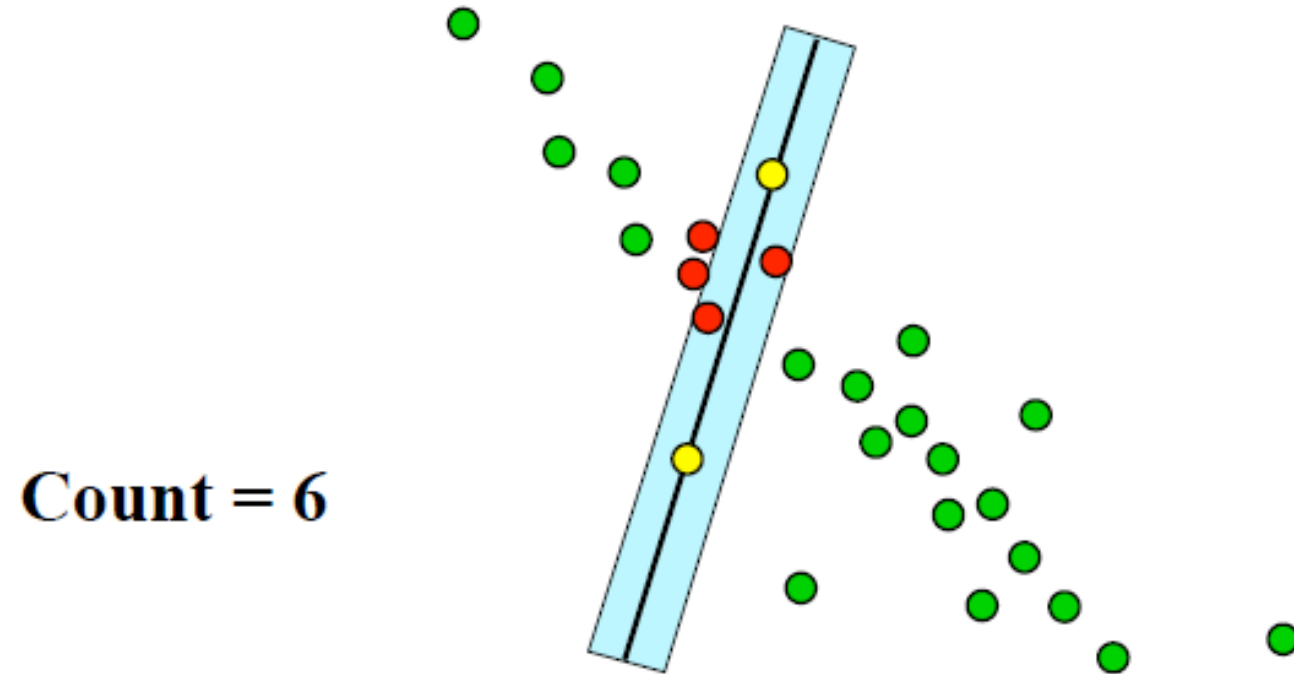
RANSAC Procedure



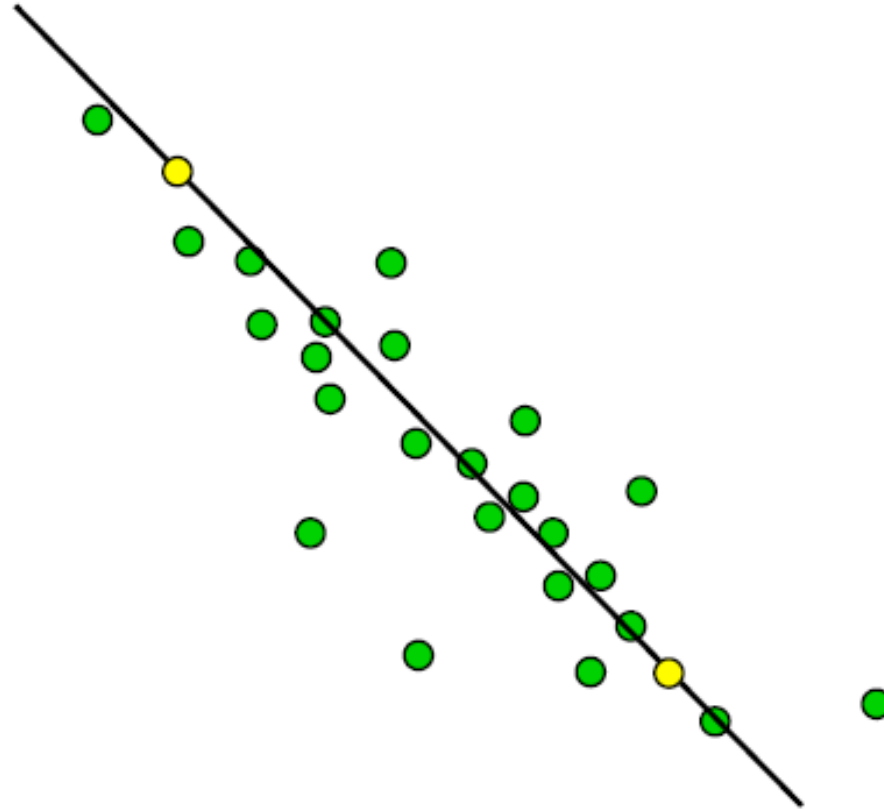
RANSAC Procedure



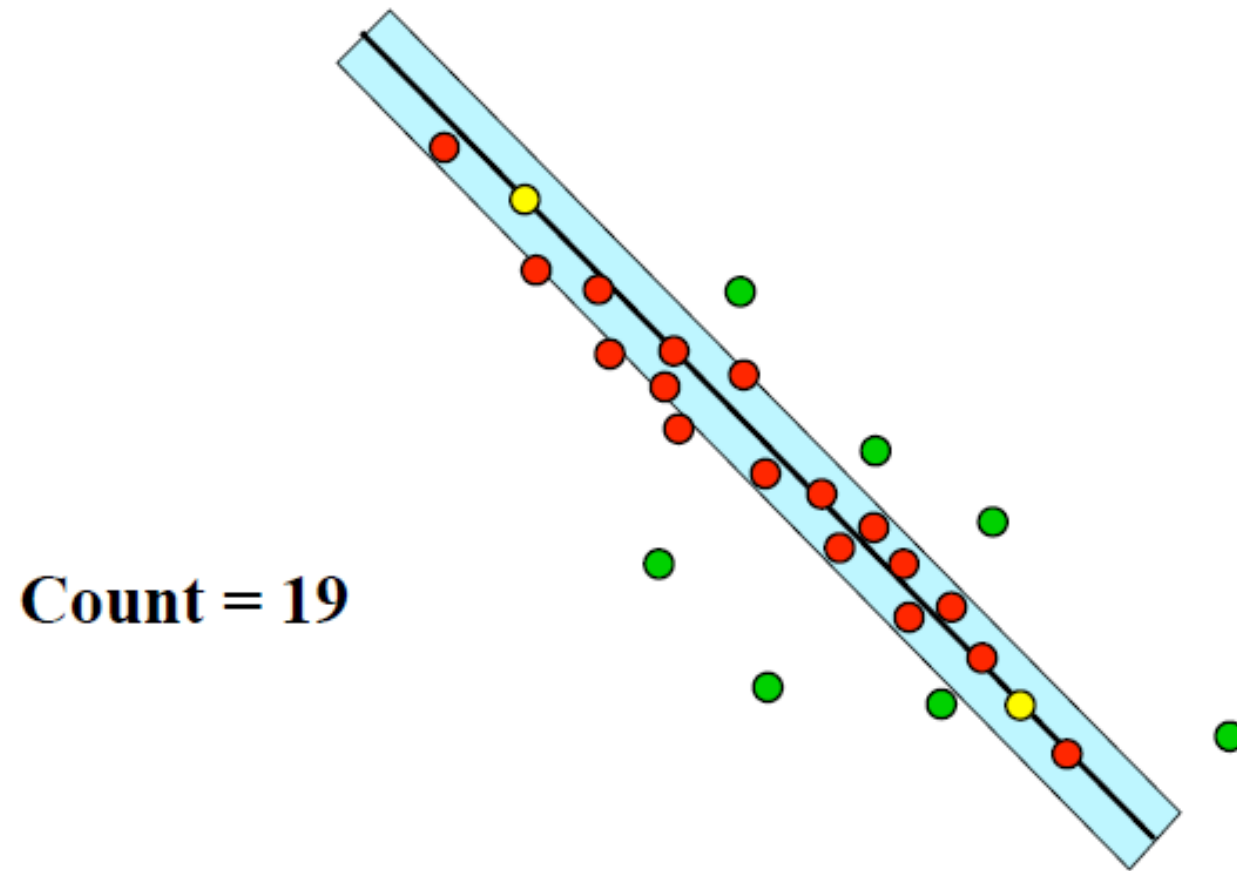
RANSAC Procedure



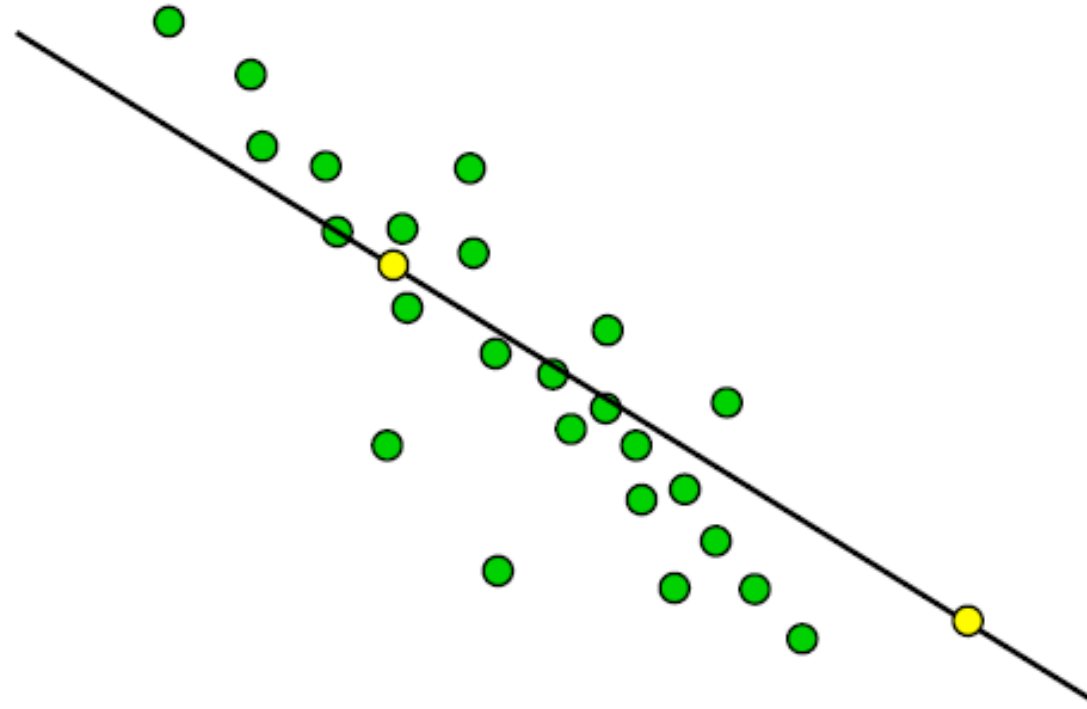
RANSAC Procedure



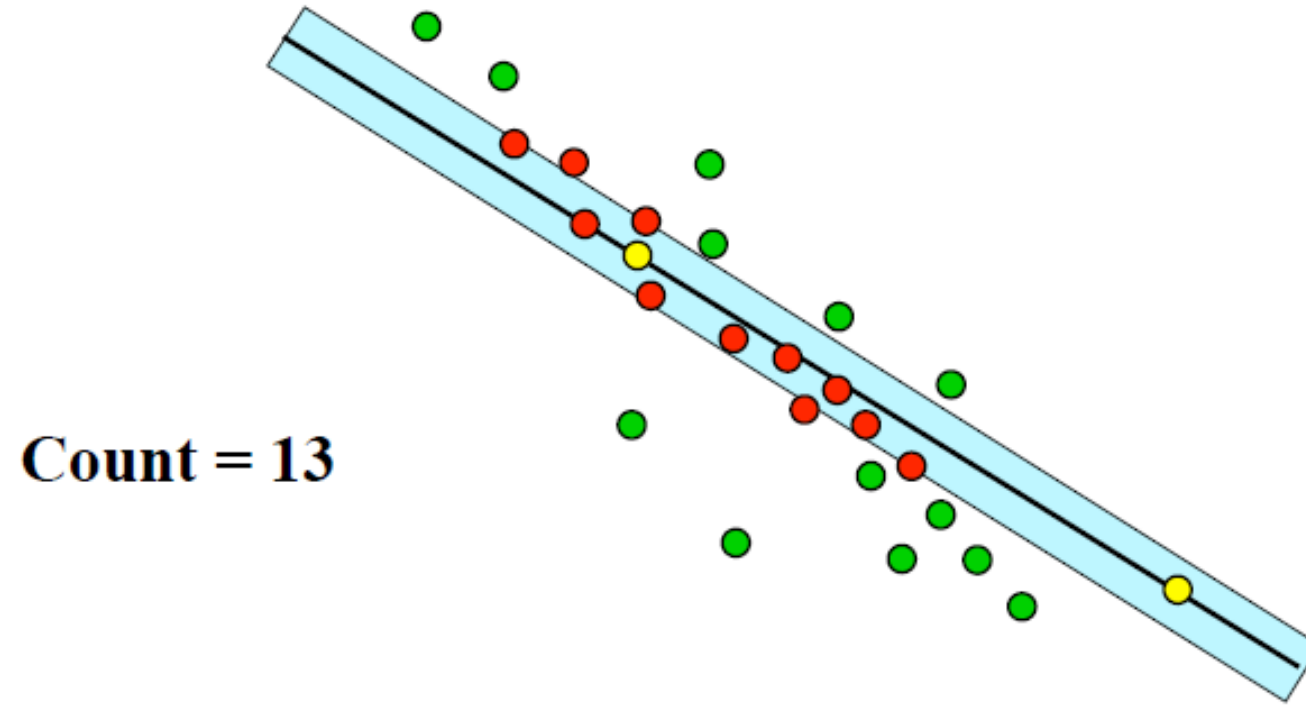
RANSAC Procedure



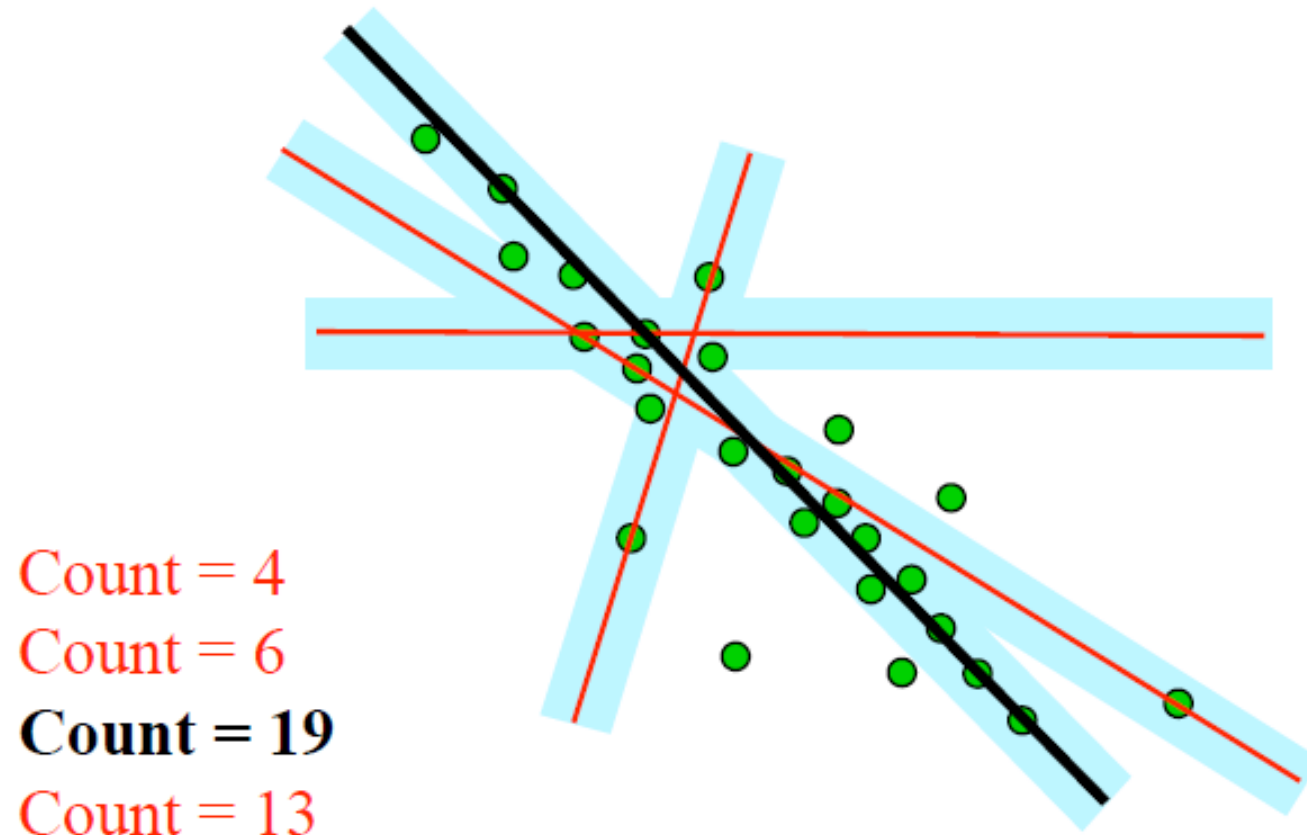
RANSAC Procedure



RANSAC Procedure



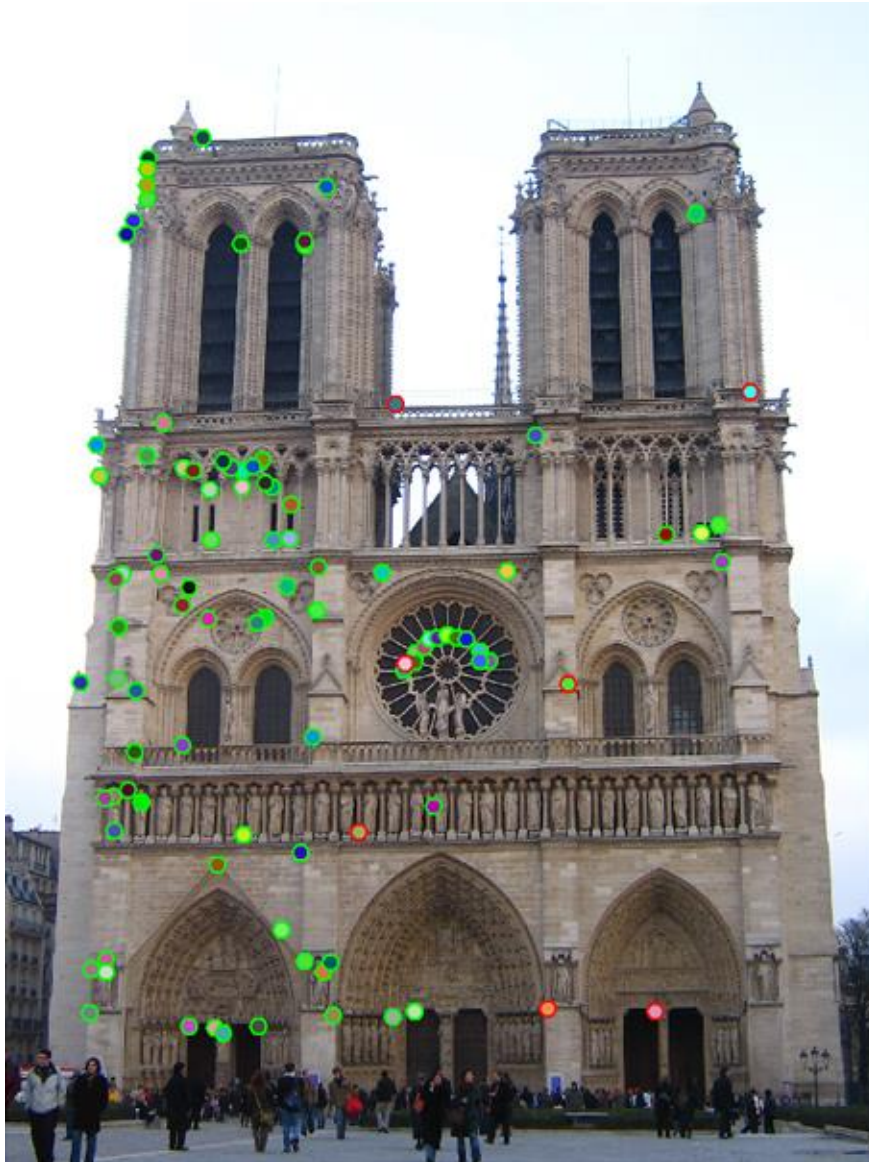
RANSAC Procedure



RANSAC - *Loop*

1. Randomly select a *seed group* of points on which to base transformation estimate (e.g., a group of matches)
2. Find *inliers* to this transformation
3. If the number of inliers is sufficiently large, re-compute least-squares estimate of transformation on all of the inliers
4. Keep the transformation with the largest number of inliers

Given matches, what Next?



Algorithm 15.4: RANSAC: fitting lines using random sample consensus

Determine:

n — the smallest number of points required

k — the number of iterations required

t — the threshold used to identify a point that fits well

d — the number of nearby points required
to assert a model fits well

Until k iterations have occurred

Draw a sample of n points from the data
uniformly and at random

Fit to that set of n points

For each data point outside the sample

Test the distance from the point to the line
against t ; if the distance from the point to the line
is less than t , the point is close

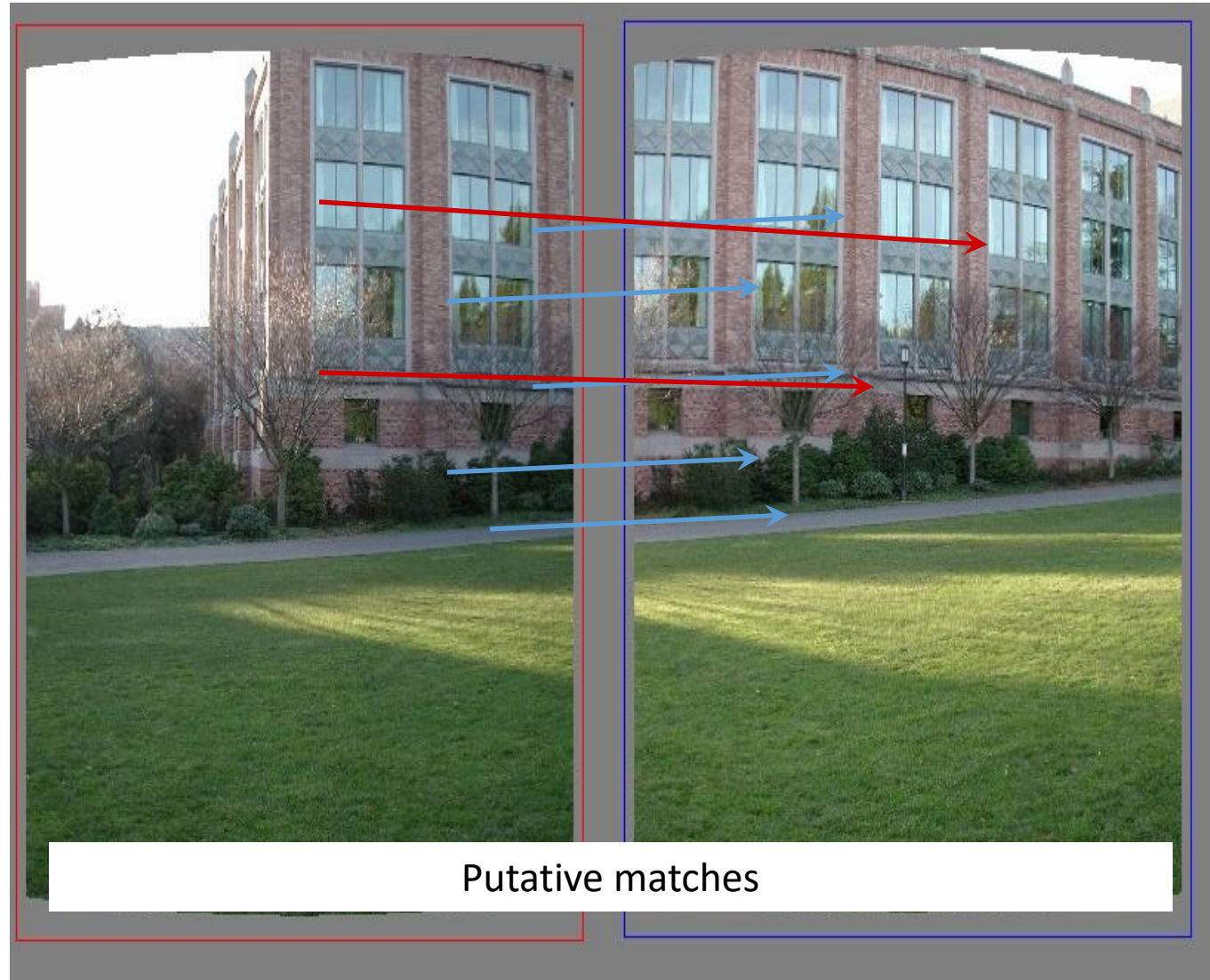
end

If there are d or more points close to the line
then there is a good fit. Refit the line using all
these points.

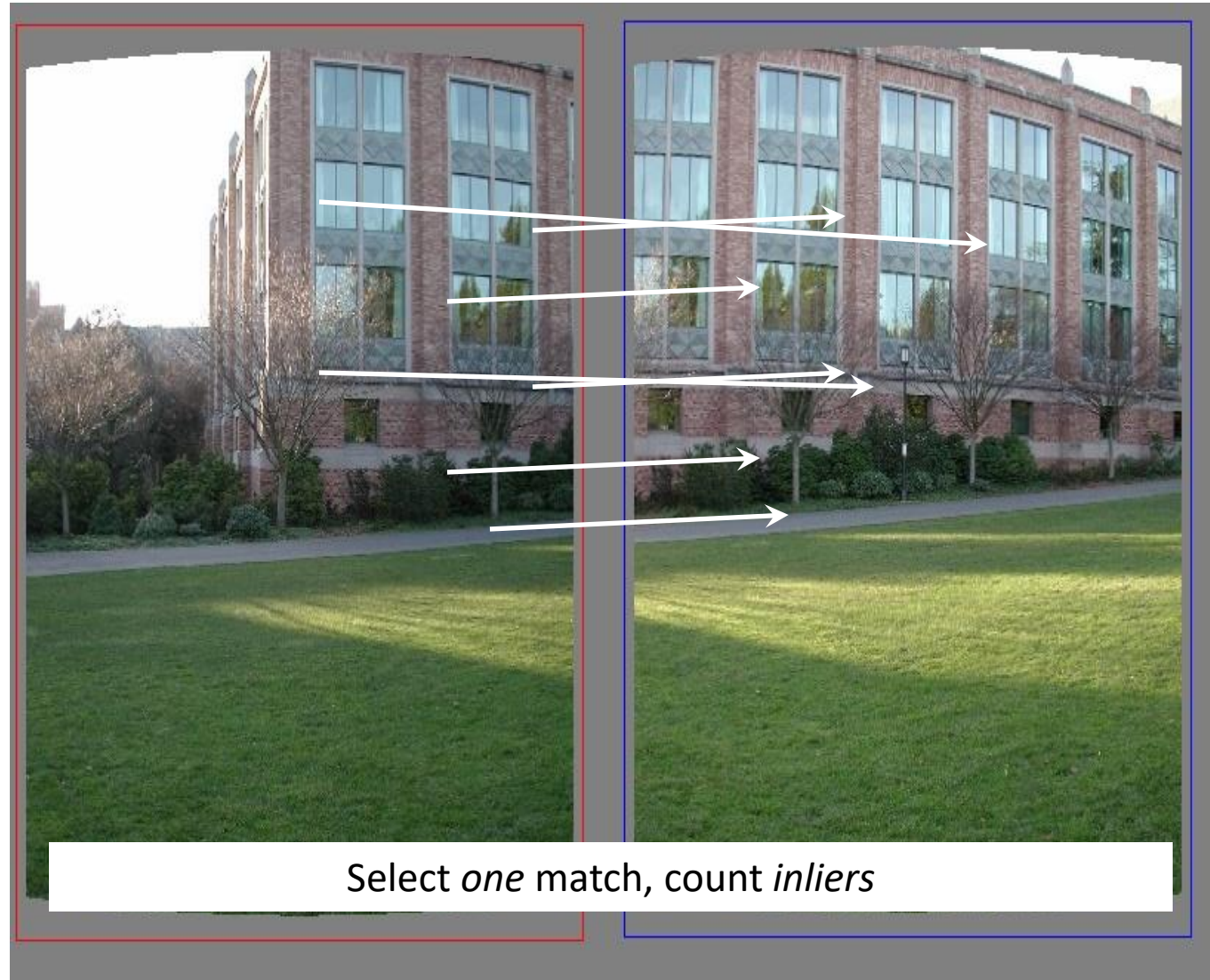
end

Use the best fit from this collection, using the
fitting error as a criterion

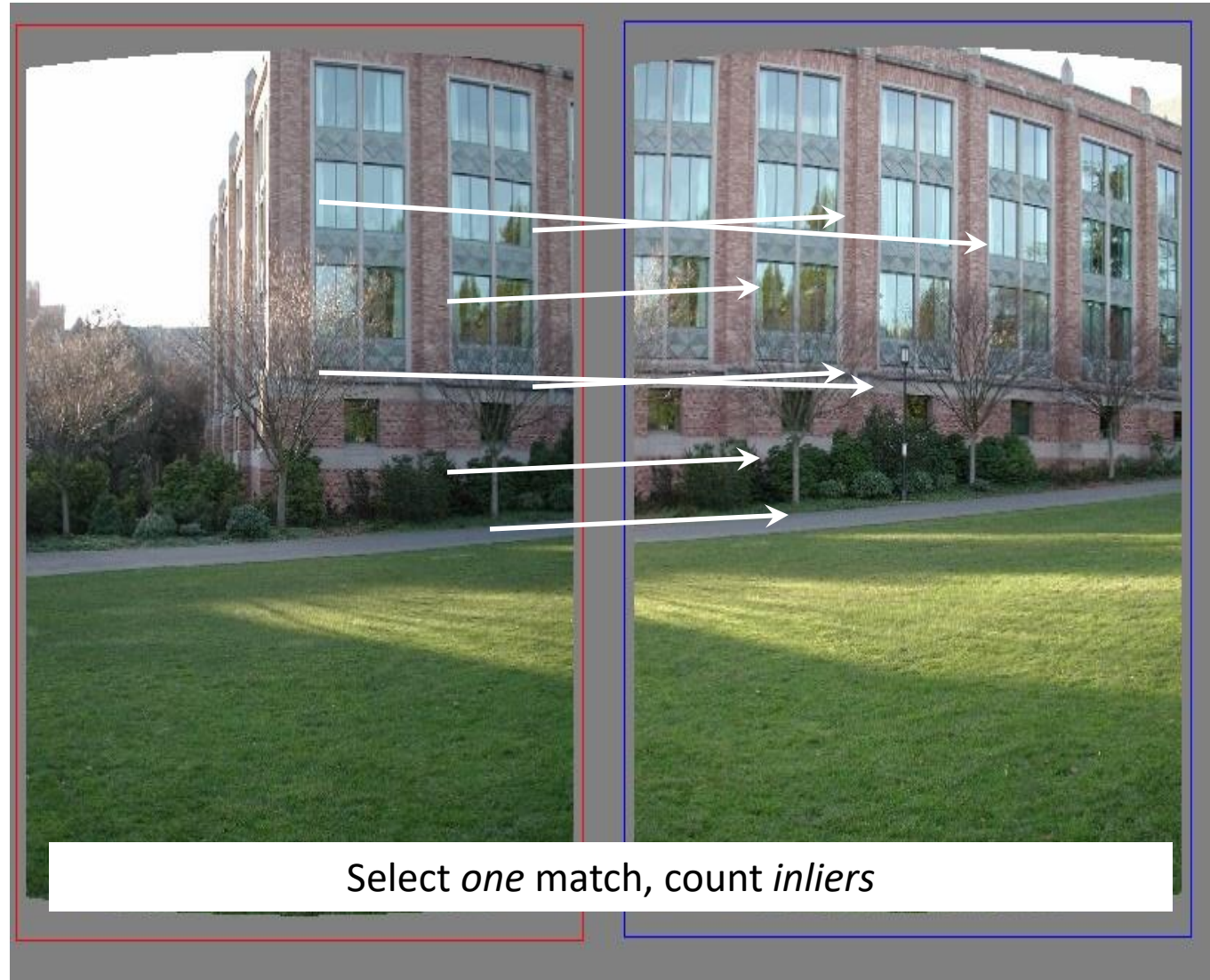
RANSAC example: Translation



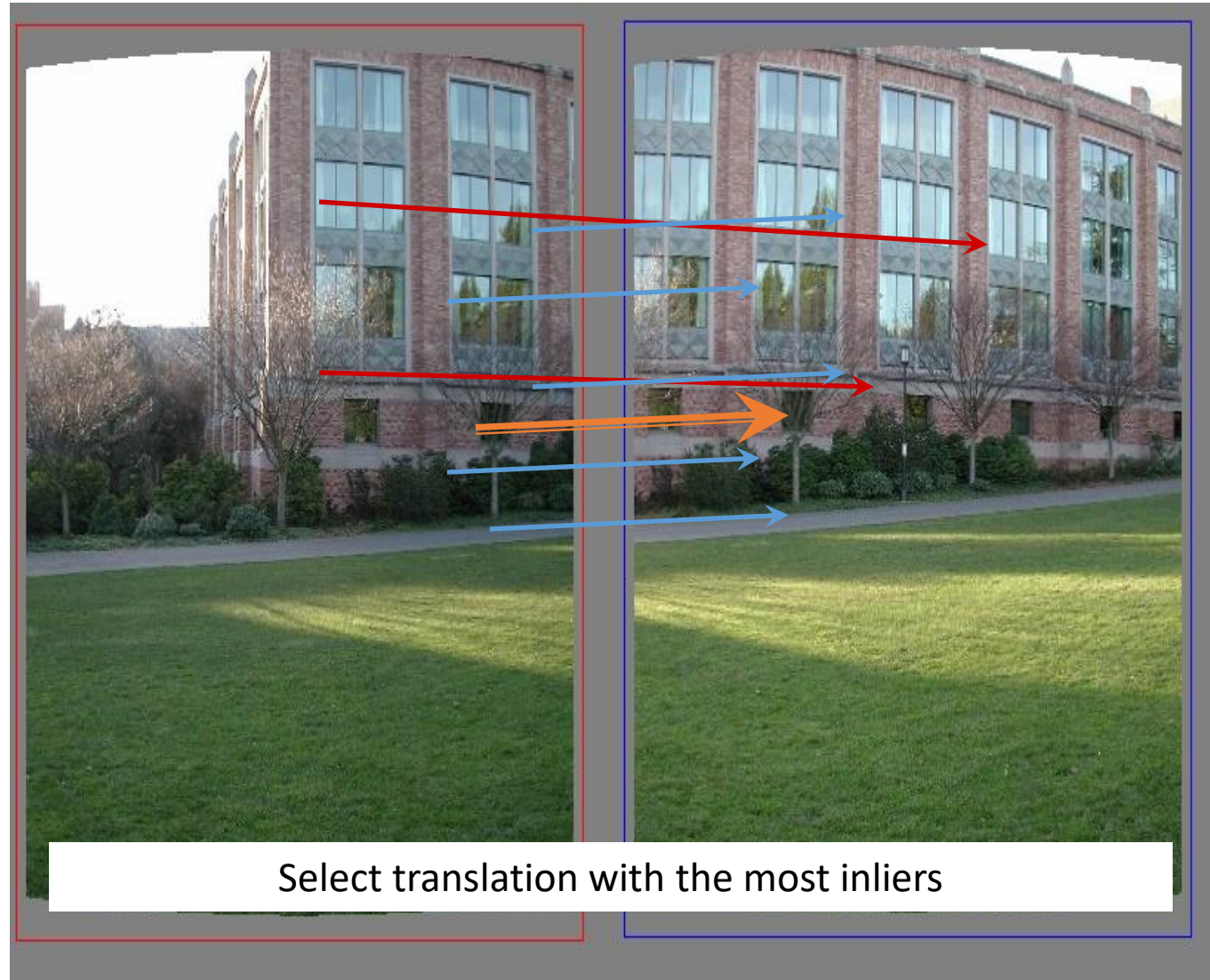
RANSAC example: Translation



RANSAC example: Translation



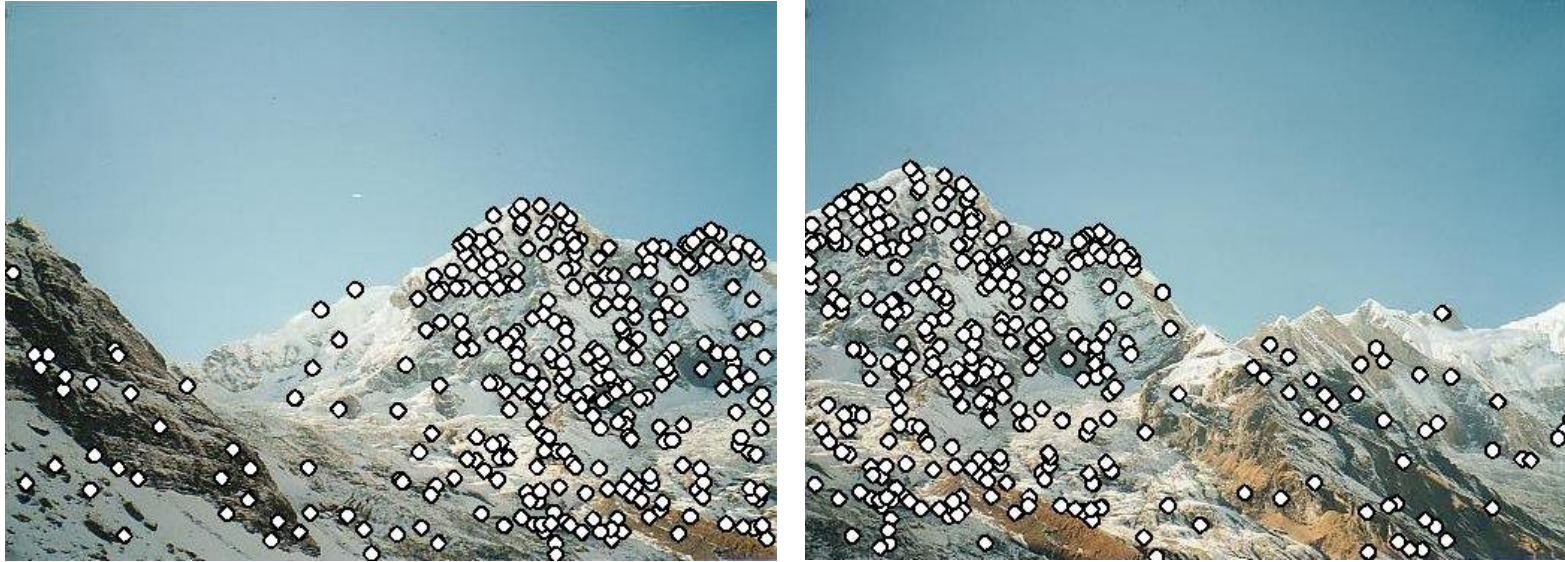
RANSAC example: Translation



Robust feature-based alignment



Robust feature-based alignment



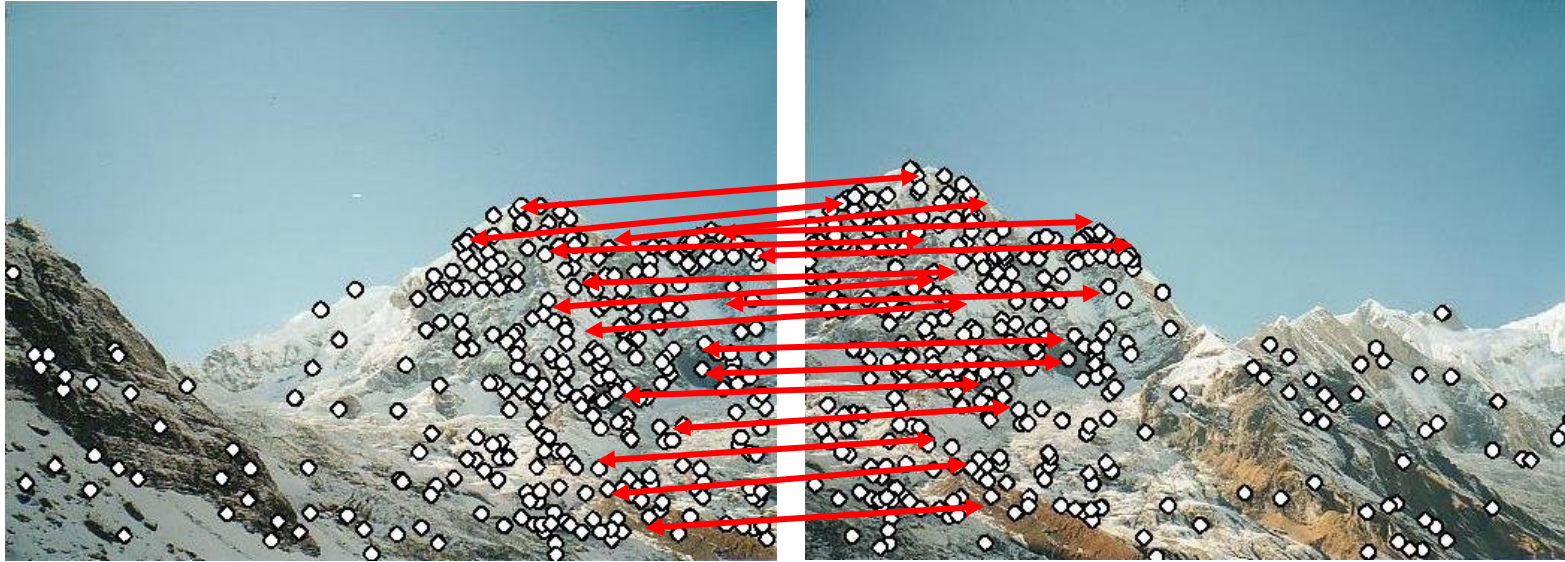
- Extract features

Robust feature-based alignment



- Extract features
- Compute *putative matches*

Robust feature-based alignment



- Extract features
- Compute *putative matches*
- Loop:
 - *Hypothesize* transformation T
 - *Verify* transformation (search for other matches consistent with T)

Robust feature-based alignment



- Extract features
- Compute *putative matches*
- Loop:
 - *Hypothesize* transformation T
 - *Verify* transformation (search for other matches consistent with T)

How Many Samples to Choose?

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?

How Many Samples to Choose?

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

$$1 - (1 - (\underbrace{1 - e}_{\text{Probability that choosing one point yields an inlier}})^s)^N = p$$

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

How Many Samples to Choose?

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

$$1 - \underbrace{(1 - (1 - e)^s)}_{\text{Probability of choosing } s \text{ inliers in a row (sample only contains inliers)}}^N = p$$

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

How Many Samples to Choose?

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

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

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

How Many Samples to Choose?

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

$$1 - \underbrace{(1 - (1 - e)^s)}_{\text{Probability that } N \text{ samples were contaminated.}}^N = p$$

Probability that N samples
were contaminated.

How Many Samples to Choose?

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

$$\underbrace{1 - (1 - (1 - e)^s)^N}_{\text{Probability that at least one sample was not contaminated}} = p$$

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. $p=0.99$

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

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

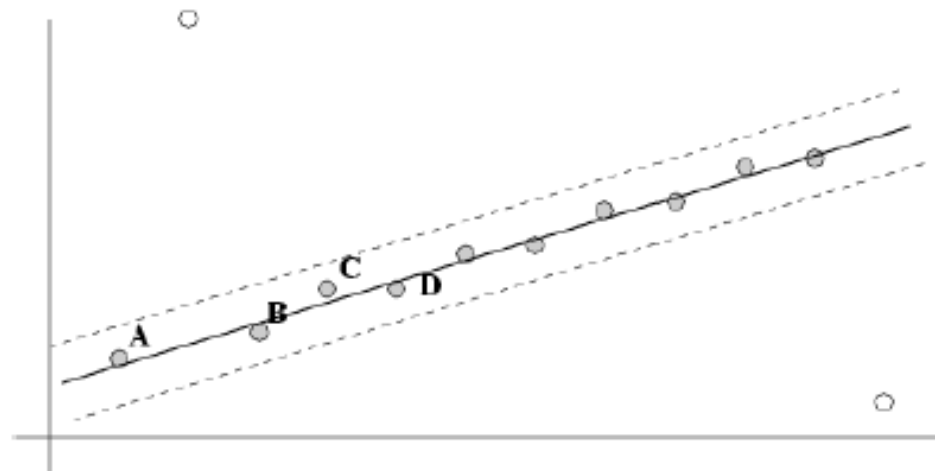
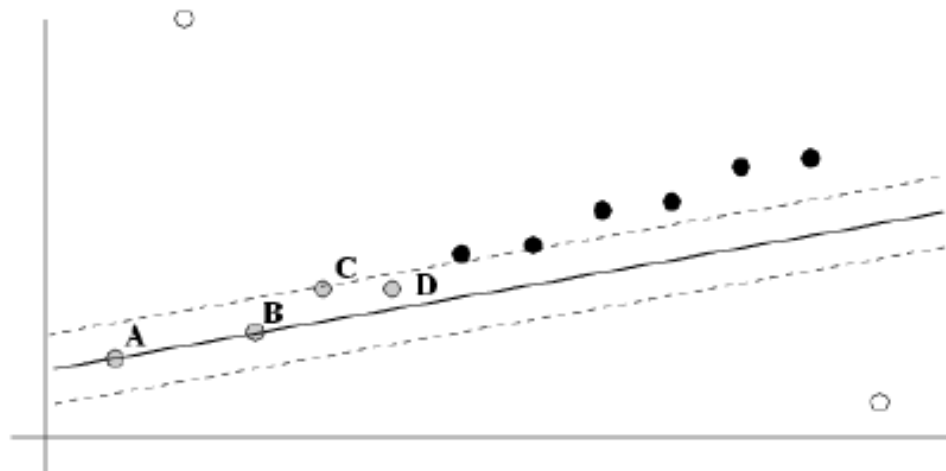
proportion of outliers e							
s	5%	10%	20%	25%	30%	40%	50%
2	2	3	5	6	7	11	17
3	3	4	7	9	11	19	35
4	3	5	9	13	17	34	72
5	4	6	12	17	26	57	146
6	4	7	16	24	37	97	293
7	4	8	20	33	54	163	588
8	5	9	26	44	78	272	1177

How to choose the parameters?

- Number of sample points S
 - Minimum number needed to fit a model
- Outlier ratio e ($e = \text{\#outliers}/\text{\#datapoints}$)
- Number of trials T
 - Choose T so that, with probability p , at least one random sample is free from outliers
- Distance threshold *delta*
 - Choose *delta* so that a good point with noise is likely (prob = 0.95) within threshold

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.



RANSAC : In Summary

Pros

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

Cons

- Computational time grows quickly with fraction of outliers and number of parameters

Common applications

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