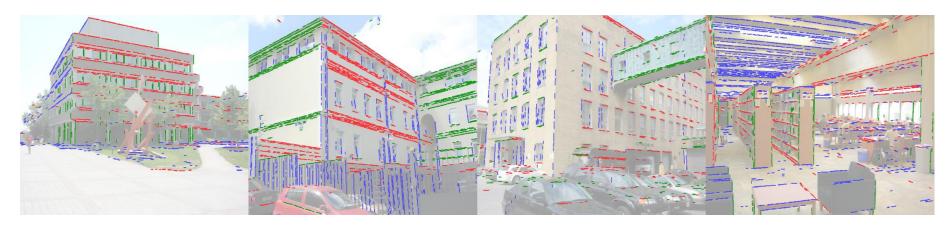
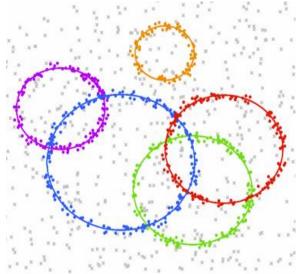


Geometric Model Fitting





with some slides stolen from Steve Seitz and Rick Szeliski



Geometric Model Fitting

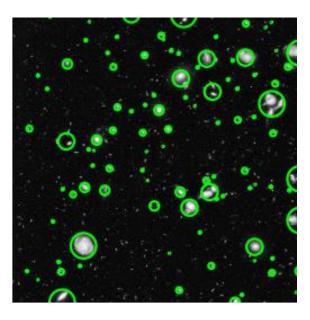
- Feature matching $(\mathbf{p}_i, \mathbf{p}'_i)$
- Model fitting (e.g. homography estimation for panoramas)
 - How many points to choose?
 - Least square model fitting
 - RANSAC (robust method for model fitting)
- Multi-model fitting problems



Flashbacks: feature detectors



Harris corners



Dog

python code from "FeaturePoints.ipynb"

from skimage.feature import corner_harris, corner_subpix, corner_peaks

hc_filter = corner_harris(image_gray)
peaks = corner_peaks(hc_filter)

from skimage.feature import blob_dog

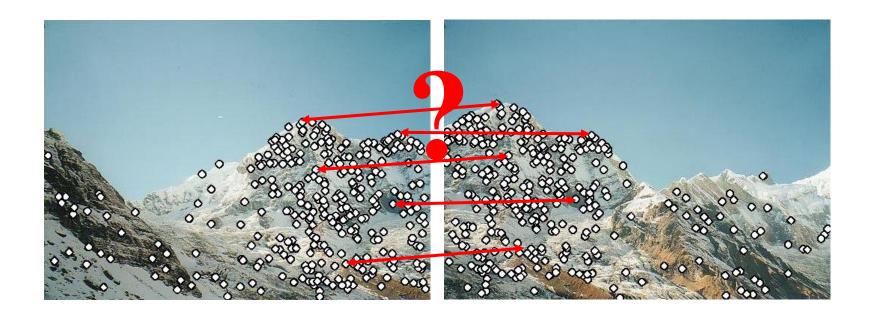
blobs = blob_dog(image_gray)



Flashbacks: feature descriptors

We know how to detect points

Next question: How to match them?



need **point descriptors** that should be

- Invariant (e.g. to gain/bias, rotation, projection, etc)
- Distinctive (to avoid false matches)

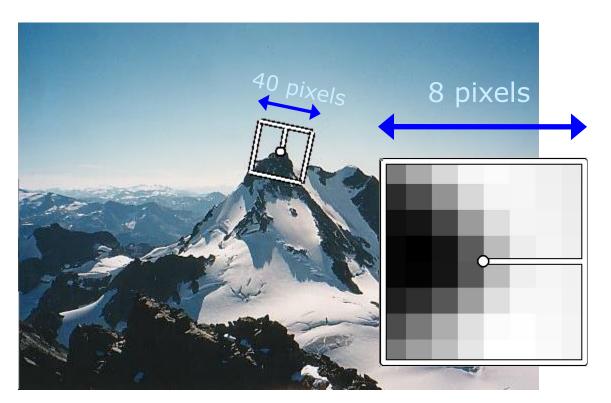


Flashbacks: MOPS descriptor

8x8 oriented patch

Sampled at 5 x scale

Bias/gain normalization: $I' = (I - \mu)/\sigma$



Another popular idea (SIFT): use gradient orientations inside the patch as a descriptor (also invariant to gain/bias)

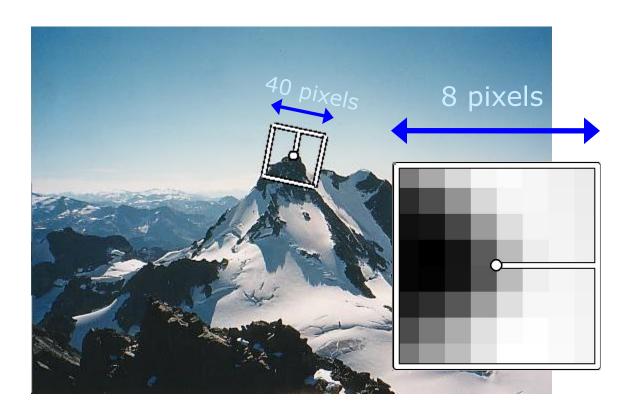


Flashbacks: MOPS descriptor

8x8 oriented patch

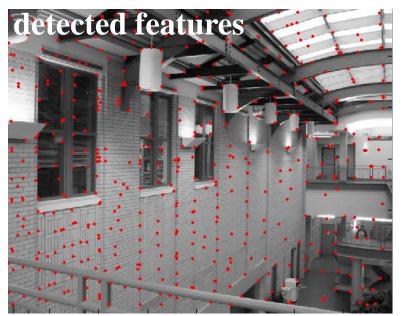
Sampled at 5 x scale

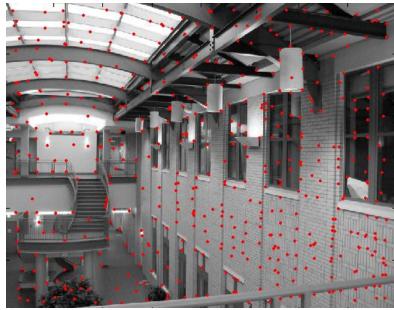
Bias/gain normalization: $I' = (I - \mu)/\sigma$

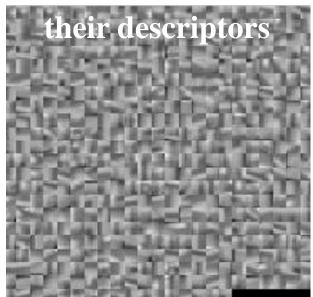


Popular descriptors: MOPS, SIFT, SURF, HOG, BRIEF, many more...

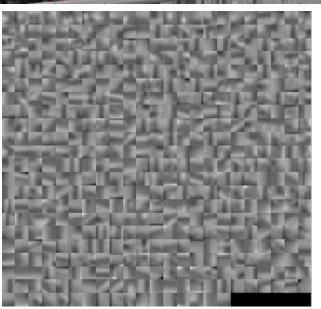














Optimal matching:

- Bipartite matching, quadratic assignment (QA) problems
 - too expensive

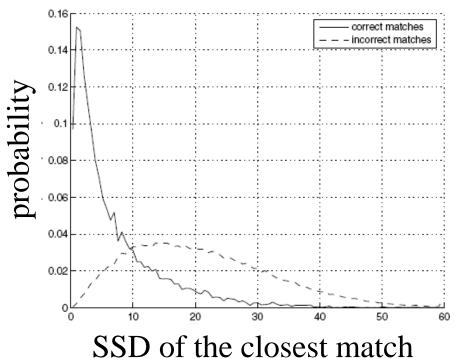
Common simple approach:

- use SSD (sum of squared differences) between two descriptors (patches).
- for each feature in image 1 find a feature in image 2 with the lowest SSD
- accept a match if SSD(patch1,patch2) < T (threshold)



SSD(patch1,patch2) < T

How to set threshold T?

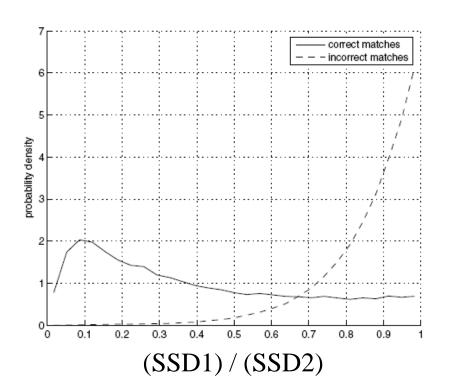


no threshold T is good for separating correct and incorrect matches



A better way [Lowe, 1999]:

- SSD of the closest match (SSD1)
- SSD of the <u>second-closest</u> match (SSD2)
- Accept the best match if it is much better than the second-best match (and the rest of the matches)

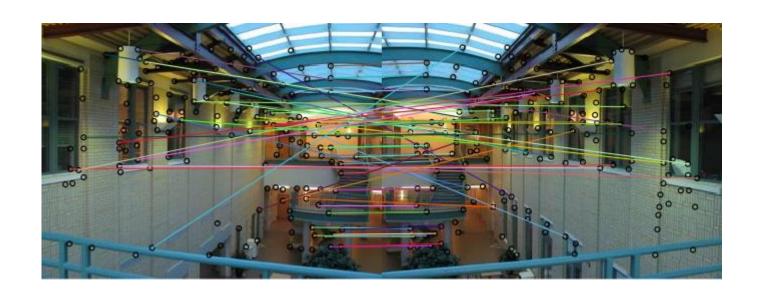


easier to select threshold T for decision test

(SSD1)/(SSD2) < T



Python example (BRIEF descriptor)



from skimage.feature import (corner_harris, corner_peaks, plot_matches, BRIEF, match_descriptors)

keypointsL = corner_peaks(corner_harris(imL), threshold_rel=0.0005, min_distance=5) keypointsR = corner_peaks(corner_harris(imR), threshold_rel=0.0005, min_distance=5)

extractor = BRIEF()

extractor.extract(imL, keypointsL) keypointsL = keypointsL[extractor.mask] descriptorsL = extractor.descriptors

extractor.extract(imR, keypointsR)
keypointsR = keypointsR[extractor.mask]
descriptorsR = extractor.descriptors

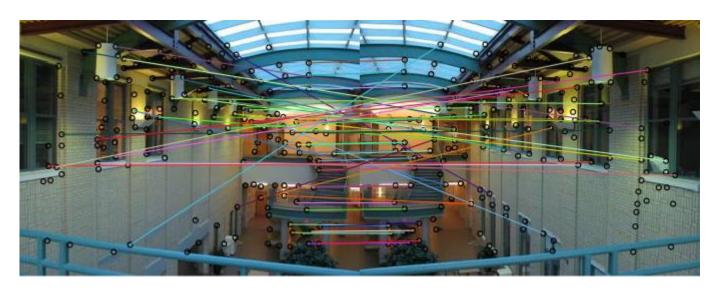
find the closest match p' for any feature p

crosscheck: keep pair (p,p') only if p is the best match for p'

matchesLR = match_descriptors(descriptorsL, descriptorsR, cross_check=True)



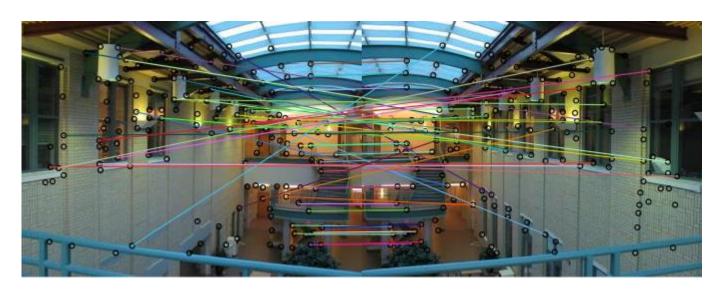
How to fit a homorgaphy???



What problems do you see for homography estimation?



How to fit a homorgaphy???



What problems do you see for homography estimation?

Issue 1: the number of matches $(\mathbf{p}_i, \mathbf{p}'_i)$ is more than 4

Answer: model fitting via "least squares" (later, slide 21)

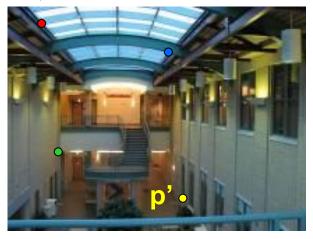
Issue 2: too many *outliers* or wrong matches $(\mathbf{p}_i, \mathbf{p}'_i)$

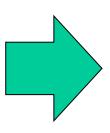
Answer: robust model fitting via RANSAC (later, slide 35)

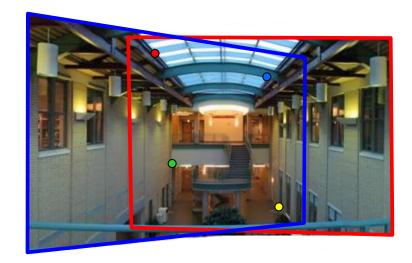


(from Topic 5)











Consider one match (point-correspondence) $p = (x, y) \rightarrow p' = (x', y')$

$$\mathbf{p'} = \mathbf{Hp}$$

$$\begin{bmatrix} wx' \\ wy' \\ w \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \\ g & h & i \end{bmatrix} \begin{bmatrix} x \\ y \\ 1 \end{bmatrix}$$

After eliminating w = gx + hy + i:

$$\Rightarrow ax+by+c-gxx'-hyx'-ix'=0$$
$$dx+ey+f-gxy'-hyy'-iy'=0$$

Two equations linear w.r.t unknown coefficients of matrix H and quadratic w.r.t. known point coordinates (x,y,x',y')



Consider 4 point-correspondences $p_i = (x_i, y_i) \rightarrow p'_i = (x'_i, y'_i)$

$$\mathbf{p'}_{i} = \mathbf{H}\mathbf{p}_{i} \qquad \begin{bmatrix} w_{i}x'_{i} \\ w_{i}y'_{i} \\ w_{i} \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \\ g & h & i \end{bmatrix} \begin{bmatrix} x_{i} \\ y_{i} \\ 1 \end{bmatrix} \qquad \text{for i=1,2,3,4}$$

$$\Rightarrow \frac{ax_i + by_i + c - gx_ix'_i - hy_ix'_i - ix'_i = 0}{dx_i + ey_i + f - gx_iy'_i - hy_iy'_i - iy'_i = 0}$$

Special case of DLT method (see p.89 in Hartley and Zisserman)

Can be written as matrix multiplication

$$\mathbf{A}_i \cdot \mathbf{h} = \mathbf{0}$$
 for i=1,2,3,4

where $\mathbf{h} = [abcdefghi]^T$ is a vector of unknown coefficients in \mathbf{H}

and \mathbf{A}_i is a 2x9 matrix based on known point coordinates x_i, y_i, x_i', y_i'



Consider 4 point-correspondences $p_i = (x_i, y_i) \rightarrow p'_i = (x'_i, y'_i)$

$$\mathbf{p'}_{i} = \mathbf{H}\mathbf{p}_{i} \qquad \Rightarrow \qquad \mathbf{A}_{i} \cdot \mathbf{h} = \mathbf{0} \qquad \text{for i=1,2,3,4}$$

$$2x9 \quad 9x1 \quad 2x1$$

or

All four matrix equations can be "stacked up" as
$$\begin{bmatrix} \mathbf{A}_1 \\ \mathbf{A}_2 \\ \mathbf{A}_3 \\ \mathbf{A}_4 \end{bmatrix} \cdot \mathbf{h} = \mathbf{0}$$
or
$$\mathbf{A} \cdot \mathbf{h} = \mathbf{0}$$

$$8x9$$

iClilcker **Q:** how many solutions for h? A: none B: one C: many



homogeneous linear equations

Recall: Homography from 4 points

Consider 4 point-correspondences $p_i = (x_i, y_i) \rightarrow p'_i = (x'_i, y'_i)$

 $\mathbf{p'}_{i} = \mathbf{H}\mathbf{p}_{i} \qquad \Rightarrow \qquad \mathbf{A} \cdot \mathbf{h} = \mathbf{0}$ for i=1,2,3,4 (*)

8 linear equations, 9 unknowns: trivial solution h=0?
All solutions h form the (right) null space of A of dimension 1,
but they represent the same transformation (as homographies can be scaled)

as discussed in topic 5,

fix norm $\|\mathbf{h}\|=1$ (later)

To find one specific solution h, for now fix one element, e.g. i = 1 this may not work more generally, should

Homography from more than 4 points

Consider *N* point-correspondences $p_i = (x_i, y_i) \rightarrow p'_i = (x'_i, y'_i)$

$$\mathbf{p'}_i = \mathbf{H}\mathbf{p}_i$$

for
$$i = 1,...,N$$

Questions:

Are there any benefits from knowing more point correspondences? What if 4 points correspondences are known with error?

over-constrained system

$$\Rightarrow \mathbf{A}_{1:8} \cdot \mathbf{h}_{1:8} = -\mathbf{A}_{9}$$
2Nx8 8x1 2Nx1

First, consider a simpler model fitting problem...



Simpler example: line fitting

Assume a set of data points $(X_1, X_1^{'})$, $(X_2, X_2^{'})$, $(X_3, X_3^{'})$, ... (e.g. person's height vs. weight)

We want to fit a linear model (a,b) to predict X' from X

$$a \cdot X + b = X'$$

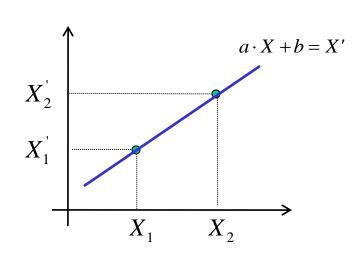
How many pairs (X_i, X_i) do we need to find a and b?

$$X_1 a + b = X_1$$
$$X_2 a + b = X_2$$

$$\begin{bmatrix} X_{1} & I \\ X_{2} & I \end{bmatrix} \begin{bmatrix} a \\ b \end{bmatrix} = \begin{bmatrix} X_{1}' \\ X_{2}' \end{bmatrix}$$

$$\mathbf{A} \cdot \mathbf{x} = \mathbf{B}$$

$$x = A^{-1} \cdot B$$





Simpler example: line fitting

Assume a set of data points $(X_1, X_1^{'})$, $(X_2, X_2^{'})$, $(X_3, X_3^{'})$, ... (e.g. person's height vs. weight)

We want to fit a linear model to predict X' from X

$$a \cdot X + b = X'$$

What if the data points (X_i, X_i) are noisy?

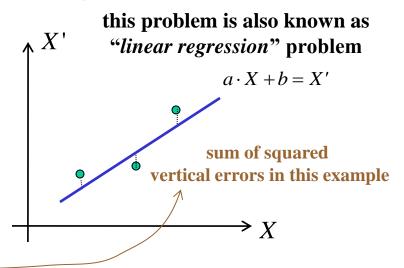
$$\begin{bmatrix} X_1 & 1 \\ X_2 & 1 \\ X_3 & 1 \end{bmatrix} \begin{bmatrix} a \\ b \end{bmatrix} = \begin{bmatrix} X_1' \\ X_2' \\ X_3' \\ \dots \end{bmatrix}$$

$$\mathbf{A} \cdot \mathbf{x} = \mathbf{B}$$

 $\min ||Ax - B||^2 \text{ (least-squares)}$



where $A^{-1} \equiv V \cdot W^{-1} \cdot U^T$ is a pseudo-inverse based on SVD decomposition $A = U \cdot W \cdot V^T$ (in python, one can use svd function in library numpy.linalg)

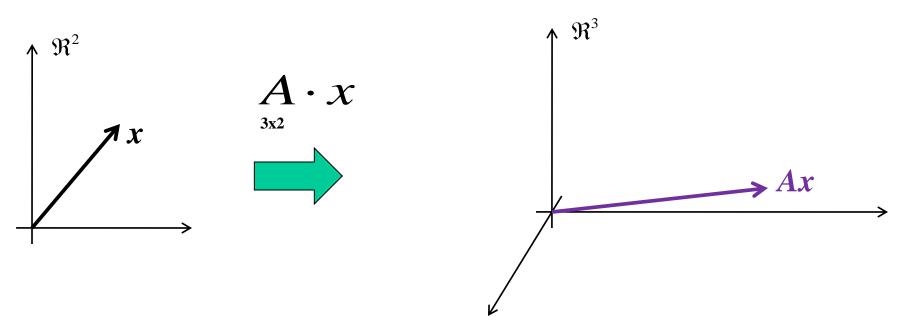




SVD: rough idea

$$A = U \cdot W \cdot V^{T}$$
Men Men Men Nen Nen Nen

where U and V are matrices with ortho-normal columns and W is diagonal with elements $w_i \ge 0$ (see "Numerical Recipes in C", edition 2, Sec. 2.6)



iClicker Moment: where are all points from \mathcal{R}^2 mapped to?

A: point

B: line

C: plane

D: whole R^3

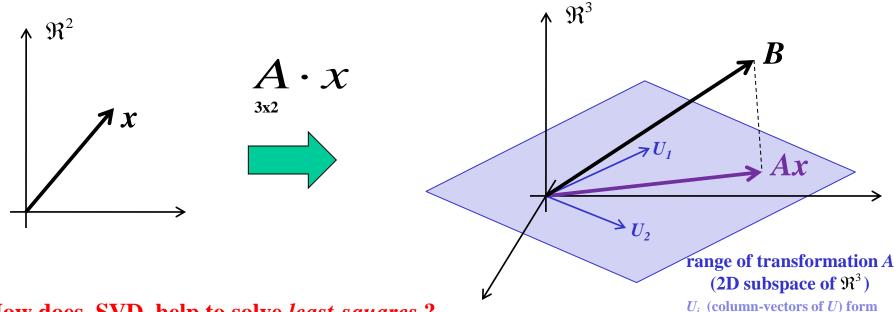


SVD: rough idea

 $M \ge N$:

$$A = \stackrel{embed}{U} \cdot \stackrel{scale}{V} \cdot \stackrel{rotate}{V}^T$$
MxN NxN NxN NxN

where U and V are matrices with ortho-normal columns and W is diagonal with elements $w_i \ge 0$ (see "Numerical Recipes in C", edition 2, Sec. 2.6)



How does SVD help to solve *least-squares*?

$$\min_{x} \|Ax - B\|^2$$

$$x = A^{-1} \cdot B$$

projection of B onto range of A

$$\equiv V \cdot W^{-1} \cdot U^T \cdot B$$

the basis of this subspace

Equivalent (fast to compute) expression

$$x = (A^T A)^{-1} \cdot A^T \cdot B$$

and: ATA - VWITWWT - VV

Indeed: $A^TA = VWU^TUWV^T = VW^2V^T$ so $(A^TA)^{-1} \cdot A^T = VW^{-2}V^T \cdot VWU^T = VW^{-1}U^T$

If M>>N computing inverse of *positive* semi-definite NxN matrix A^TA can be faster than SVD of MxN matrix A



Least squares line fitting

Data generated as $X'_i = a X_i + b + \delta X_i$ for Normal noise δX_i



$$\min_{x=(a,b)^{\top}} \|Ax - B\|^2 \xrightarrow{\qquad \qquad } x = A^{-1}B$$
 least squares



Homography from N ≥ 4 points

Consider N point correspondences $p_i = (x_{i,} y_i) \rightarrow p'_i = (x'_{i,} y'_i)$

$$\mathbf{p'}_{i} = \mathbf{H}\mathbf{p}_{i}$$
 \Rightarrow $\mathbf{A} \cdot \mathbf{h} = \mathbf{0}$ (*)

for $i = 1,...,N$ over-constrained system

Approach 1: add constraint i = 1. So, there are only 8 unknowns.

Set up a system of linear equations for vector of unknowns $h_{1:8} = [a,b,c,d,e,f,g,h]^T$

$$\mathbf{A}_{1:8} \cdot \mathbf{h}_{1:8} = \mathbf{B} = -\mathbf{A}_{9}$$
_{2Nx8} 8x1 2Nx1

$$\min_{h_{1:8}} ||A_{1:8}h_{1:8} - B||^2$$
 (least-squares)

compute inverse for $\mathbf{A}_{1:8}^T \mathbf{A}_{1:8}$ as in line fitting, then $\mathbf{h}_{1:8} = (\mathbf{A}_{1:8}^T \cdot \mathbf{A}_{1:8})^{-1} \cdot \mathbf{A}_{1:8}^T \cdot (-\mathbf{A}_9)$



Homography from N ≥ 4 points

Consider N point correspondences $p_i = (x_{i,} y_i) \rightarrow p'_i = (x'_{i,} y'_i)$

$$\mathbf{p'}_{i} = \mathbf{H}\mathbf{p}_{i}$$
 \Rightarrow $\mathbf{A} \cdot \mathbf{h} = \mathbf{0}$ (*)

for $i = 1,...,N$ over-constrained system

Approach 2: add constraint $\|\mathbf{h}\|=1$

solve
$$\min_{h:||h||=1} ||A\mathbf{h}||^2$$
 (homogeneous least-squares)

Solution: (unit) eigenvector of $\mathbf{A}^T \mathbf{A}$ corresponding to the smallest eigen-value (use SVD, see next slide)

DLT method (see p.91 in Hartley and Zisserman)



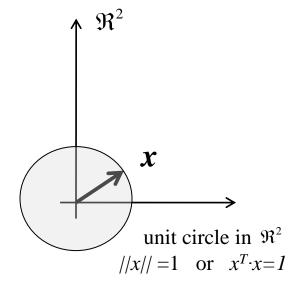
Simple motivating example:

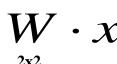
Consider 2x2 diagonal matrix

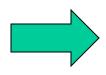
$$W = \begin{vmatrix} w_1 & 0 \\ 0 & w_2 \end{vmatrix}$$

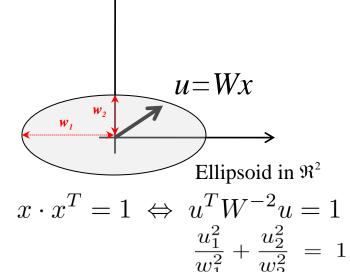
solve:

$$\min_{x:||x||=1}\|W\cdot x\|$$









Solution: x = (1,0) if $w_1 < w_2$ x = (0,1) if $w_2 < w_1$

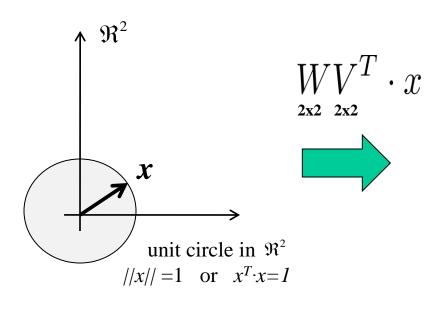
 \leftarrow equivalently, solve $\min_{u \in Ellips} || u ||$

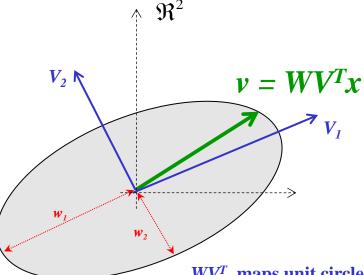


General case: use SVD (rough idea)

$$A = \stackrel{\it embed}{U} \cdot \stackrel{\it scale}{V} \cdot \stackrel{\it rotate}{V}^T$$

where U and V are matrices with ortho-normal columns and W is diagonal with elements $w_i \ge 0$ (see "Numerical Recipes in C", edition 2, Sec. 2.6)





can interpret multiplication by V^T as change of ortho-normal basis (or as space rotation, but illustration should be modified)

• interpret multiplication by W as anisotropic scaling/deformation

 WV^T maps unit circle $x^T \cdot x = 1$ onto ellipsoid $v^T W^{-2} v = 1$ in basis $\{V_1, V_2\}$

check that $x=V_i$ is mapped to point $W V^T x = w_i e_i$ vector with zeros

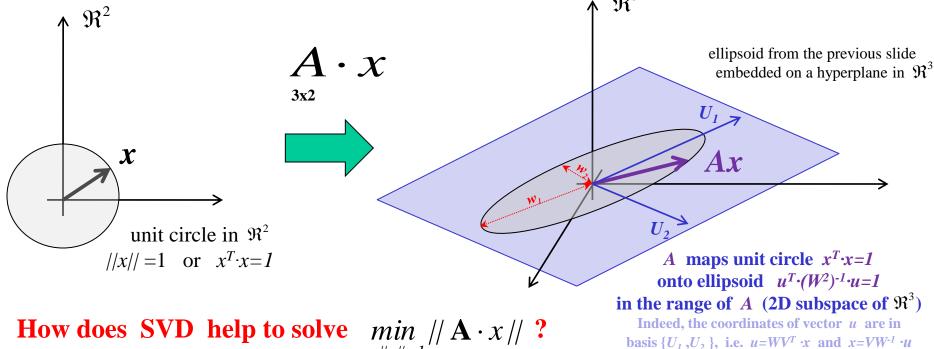
and 1 in the *i*-th position



General case: use SVD (rough idea)

$$A = U \cdot W \cdot V^T$$
MxN MxN NxN NxN NxN

where U and V are matrices with ortho-normal columns and W is diagonal with elements $w_i \ge 0$ (see "Numerical Recipes in C", edition 2, Sec. 2.6)



How does SVD help to solve $min /| \mathbf{A} \cdot x /|$?

M > N:

$$\min_{x:||x||=1}$$
 | $\mathbf{A} \cdot x$ | ?

 $A \cdot V_i = UWV^T \cdot V_i = UW \cdot e_i = w_i \cdot (U \cdot e_i) = w_i \cdot U_i$ \Rightarrow $//A \cdot V_i / \models w_i$ vector $x = V_i$ corresponding to the least w_i solves the problem

Check that V_i and $(w_i)^2$ are eigen vectors/values for matrix A^TA (note: ellipsoid $x^T \cdot (A^TA) \cdot x = 1$ maps onto circle $u^T \cdot u = 1$)

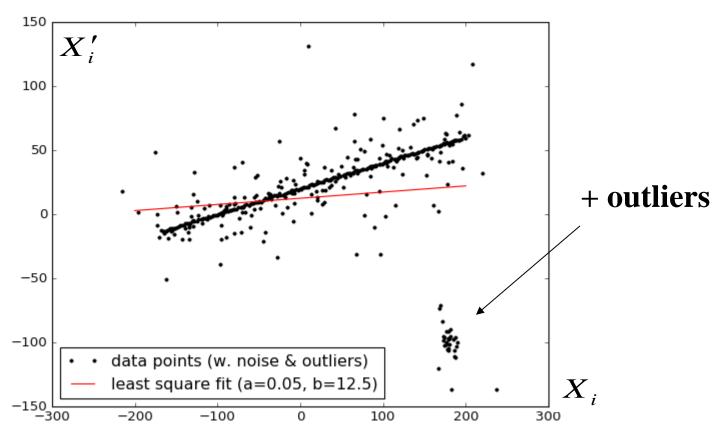
$$A^T A \cdot V_i = VW^2 V^T \cdot V_i = VW^2 \cdot e_i = w_i^2 \cdot V_i$$
 => can use eigen decomposition of $A^T A$ instead of SVD of A .

Least squares

fail in presence of outliers



Data generated as $X'_i = a X_i + b + \delta X_i$ for Normal noise δX_i



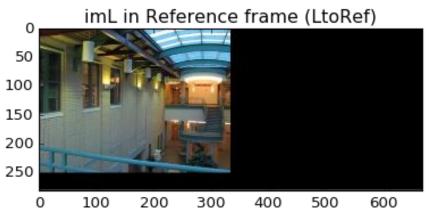
$$\min_{x=(a,b)^{\top}} \|Ax - B\|^2 \longrightarrow x = A^{-1}B$$
least squares

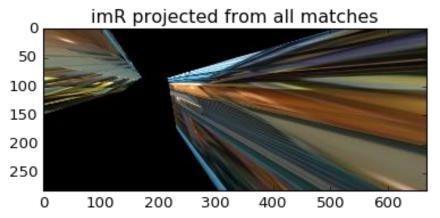
Least squares

fail in presence of outliers



 $\mathbf{h} = V_i$





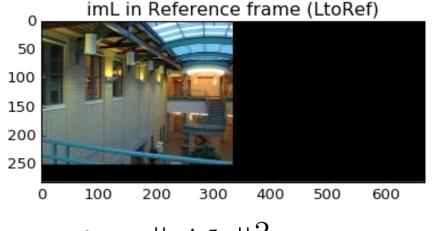
for $A = UWV^{\top}$ and $i = \arg\min w_i$

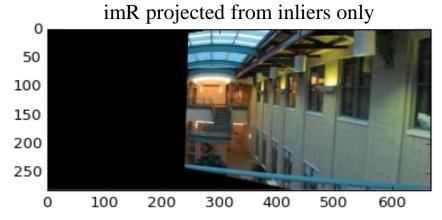
Least squares work if using "inliers" only



(detecting these? – soon)







 $\mathbf{h} = V_i$ for $A = UWV^{\top}$ and $i = \arg\min w_i$

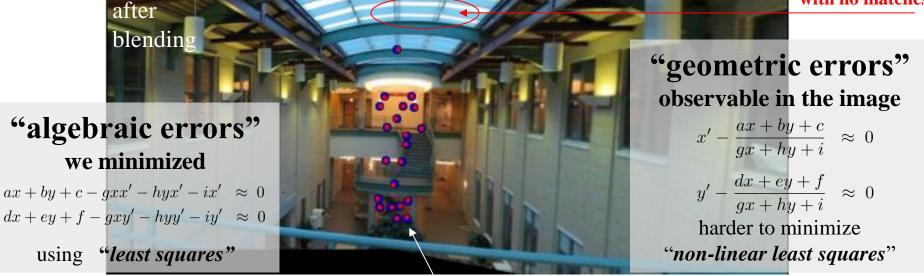
Least squares work if using "inliers" only



(detecting these? – soon)



larger errors in the area with no matches



"errors" among inliers

 $\|p'-Hp\|$

Did we actually minimize these errors?



(detecting these? – soon)





Question: how to remove outliers automatically?



Model fitting robust to outliers

We need a method that can separate inliers from outlliers

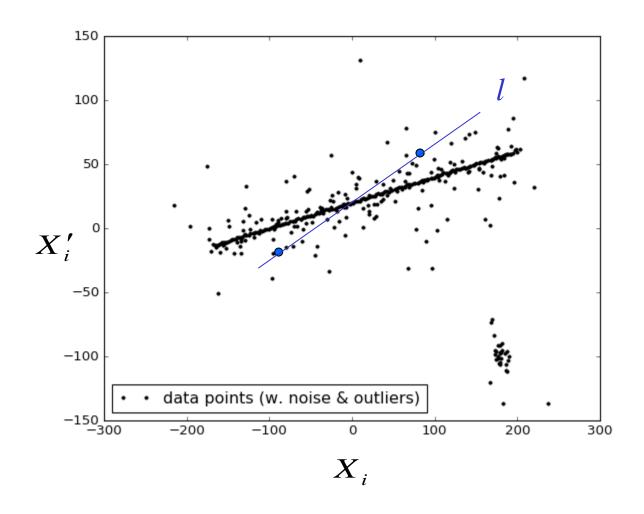
RANSAC

random sampling consensus

[Fischler and Bolles, 1981]

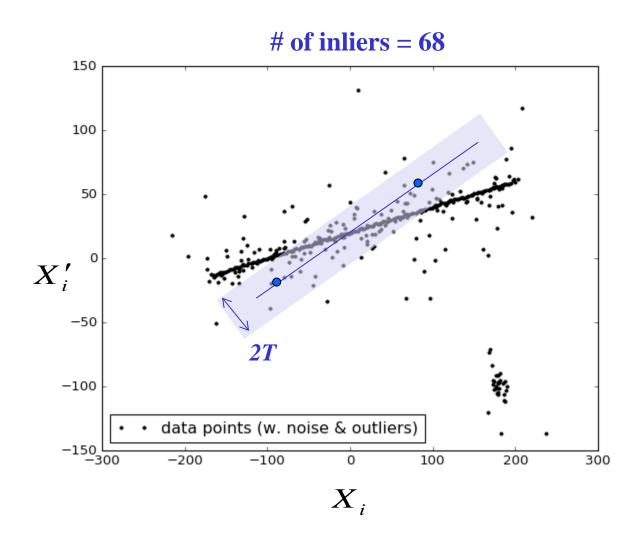


RANSAC (for line fitting example)



1. randomly sample two points from the set, get a line

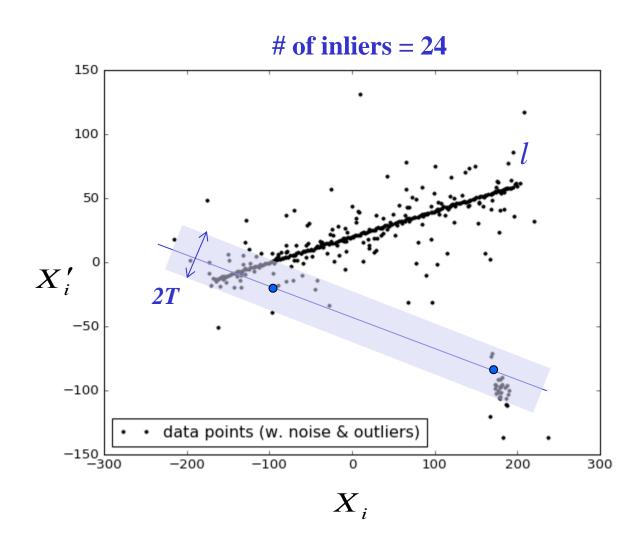




- 1. randomly sample two points from the set, get a line
 - 2. count inliers *p* for threshold *T*

$$\|p-l\| \le T$$



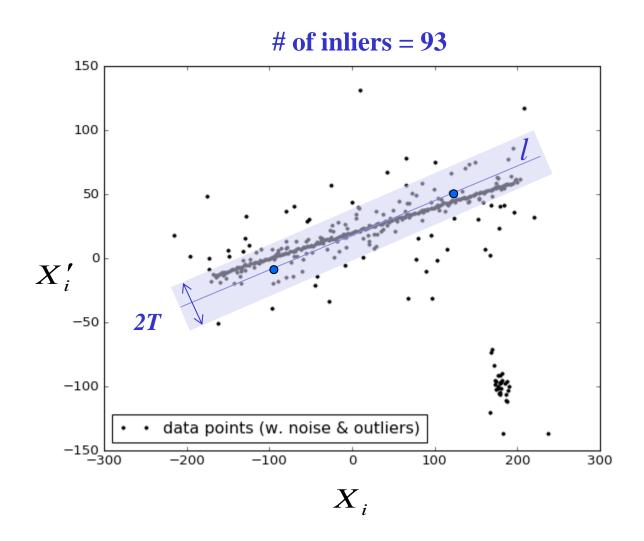


- 1. randomly sample two points from the set, get a line
 - 2. count inliers *p* for threshold *T*

$$\|p-l\| \le T$$

3. repeat



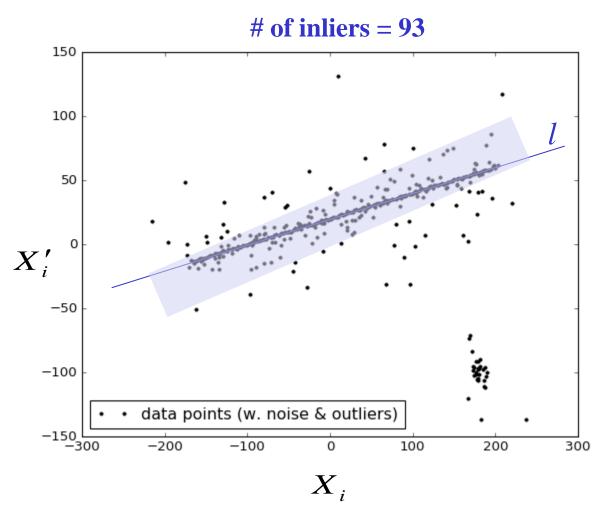


- 1. randomly sample two points from the set, get a line
 - 2. count inliers *p* for threshold *T*

$$\|p-l\| \le T$$

3. repeat N times and select model with most inliers





- 1. randomly sample two points from the set, get a line
 - 2. count inliers *p* for threshold *T*

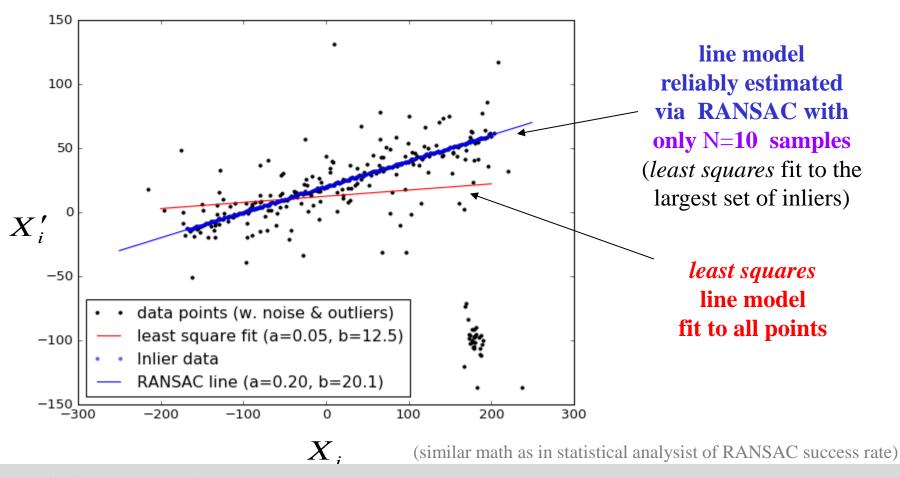
$$\|p-l\| \le T$$

- 3. repeat N times and select model with most inliers
- 4. Use *least squares* to fit a model (line) to this largest set of inliers

Q: Assume know percentage of outliers in the data.

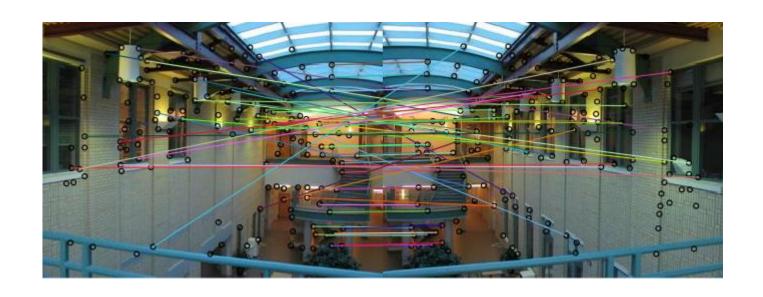
How many pairs of points (N) should be sampled to have high confidence (e.g. 95%) that at least one pair consist of two inliers? [Fischler and Bolles, 1981]





Birthday Paradox: in a group of random 23 people the probability that at least two have same birthday is 50.7%





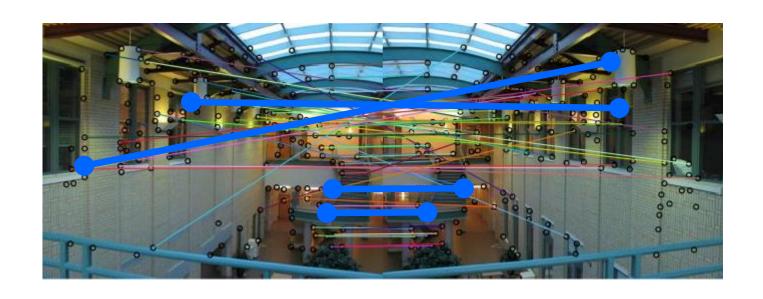
only two differences: 1. need to randomly sample four pairs (p, p')

the minimum number of matches to estimate a homography H

2. pair (p, p') counts as an inlier for a given homography H if

$$||p'-Hp|| \leq T$$





Homography for corrupted four matches is likely to have only a few inliers (p, p')

$$||p'-Hp|| \leq T$$

(randomly sampled)





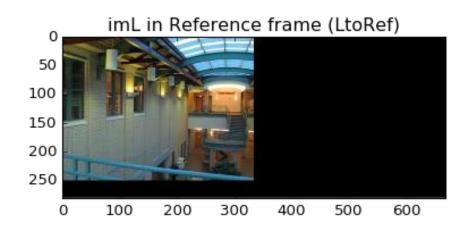
Homography for good four matches has 21 inliers (p, p')

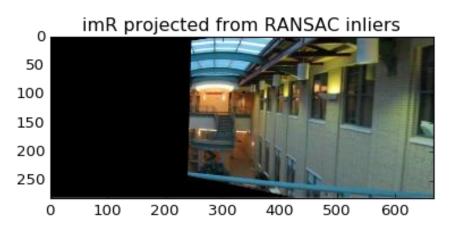
(randomly sampled)





Inliers for the randomly sampled homography with the largest inlier set











matched inliers

||p'-Hp||

The final automatic panorama result



RANSAC for robust model fitting

In general (for other models):

always sample the smallest number
of points/matches needed to estimate a model

RANSAC loop:

- Select <u>four</u> feature pairs (at random)
- 2. Compute homography H (exact)
- 3. Count *inliers* (p, p') where $||p'-Hp|| \le T$
- 4. Iterate N times (steps 1-3). Keep the largest set of inliers.
- 5. Re-compute <u>least-squares</u> H estimate on all of the inliers

 e.g. for algebraic errors

 (for simplicity)



images from <u>different</u> view points (optical centers)





Merton College III data from Oxford's Visual Geometry Group

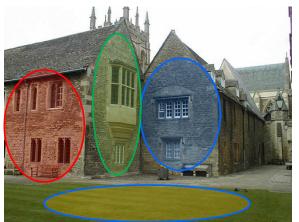
http://www.robots.ox.ac.uk/~vgg/data/data-mview.html

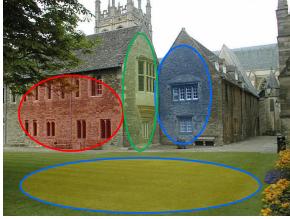
Question: Is it possible to create a panorama from these images? (or, is there a *homography* that can match overlap in these images?)

Can a *homography* map/warp **a part** of the left image onto **a part** of the right image?



images from <u>different</u> view points (optical centers)





Merton College III data

from Oxford's Visual Geometry Group

http://www.robots.ox.ac.uk/~vgg/data/data-mview.html

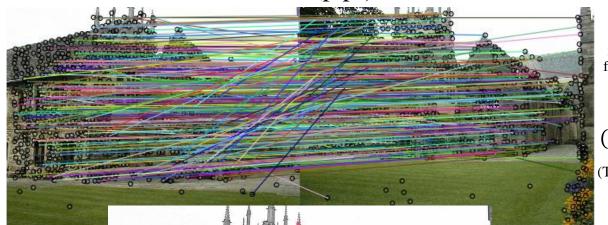
There should be a *homography* for each plane in the scene (Why?)

Question: How can we detect such *homographies*?

What do these multiple *homographies* give us?







NOTE:

good matches can be used for reconstructing 3D points if camera positions & orientations are known:

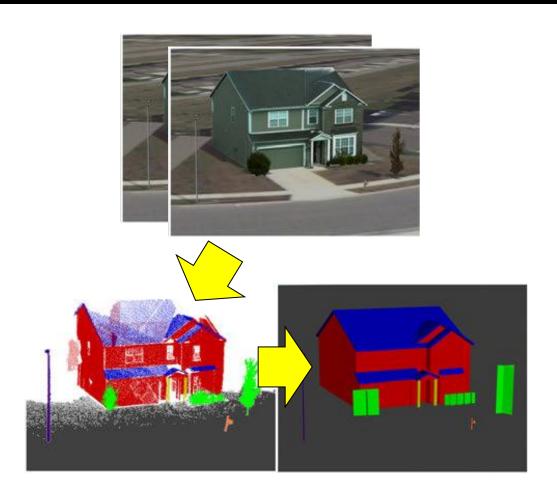
$$(p,p') \rightarrow X_p \in R^3$$

(Triangulation, see next topic)

- 1. Allow to remove bad matches (outliers)
- 2. Correspondences can be estimated with subpixel precision $(p, p') \rightarrow (p, Hp)$
- [Isack, et al. IJCV12]
- 3. Inliers allow to segment planar regions
- 4. Segmentation allows to extend correspondence from inliers to any point *p* inside segment: (*p*, *Hp*)
- 5. Piece-wise planar3D scene reconstruction

What do these multiple homographies give us?



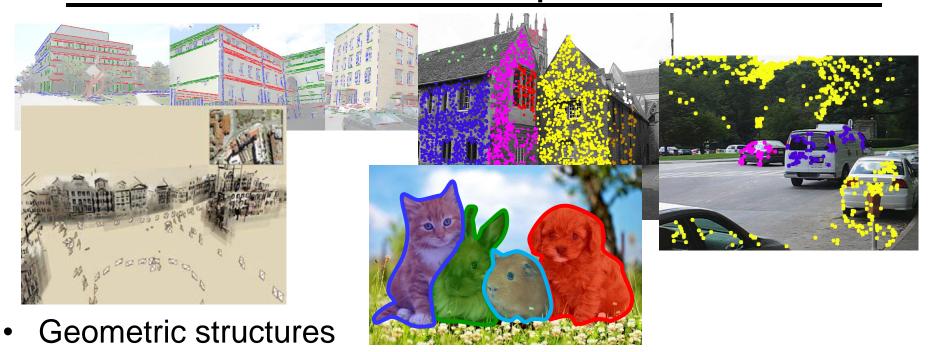


could be used for

3D piecewise planar reconstruction
from multiple views



Other K-model estimation problems in vision



- lines, planes, rigid motions (fundamental matrices), flex. models (pose)
- Stereo or multi-view reconstruction

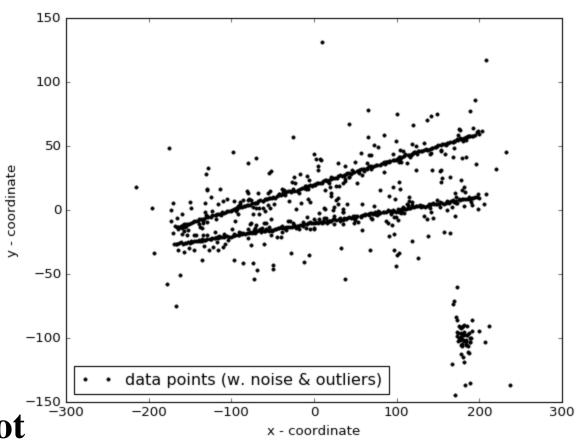
Topics 7,8,12

- projection matrices (camera positions), points and surfaces in 3D
- Object models (segmentation, classification) Topics 9,10,11,12
 - appearance models, boundary surface models, (linear & non-linear) discriminative models
 Computer vision can answer what and where

based on "model" estimation

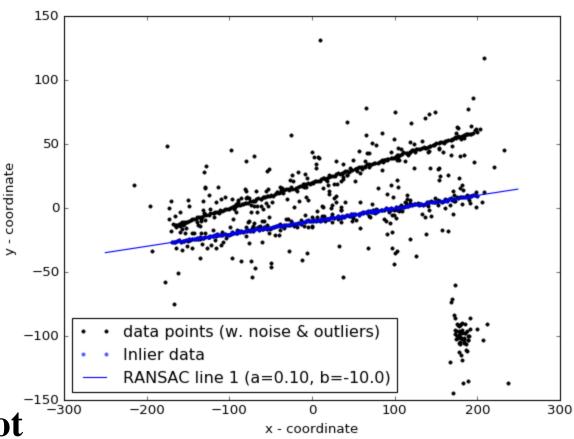
model formulation + loss function (data & regularization) + optimization

So, how do we fit multiple geometric models?



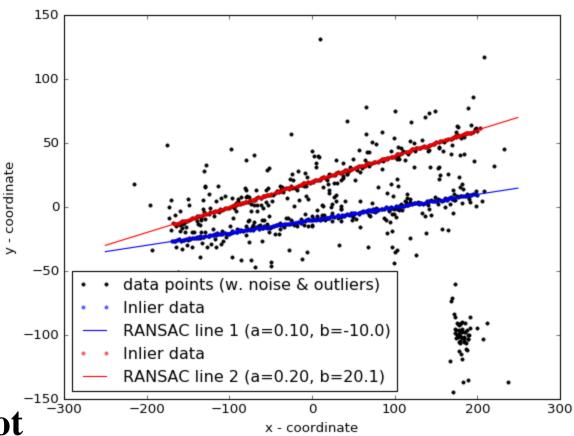
Why not RANSAC again?

So, how do we fit multiple geometric models?



Why not RANSAC again?

So, how do we fit multiple geometric models?

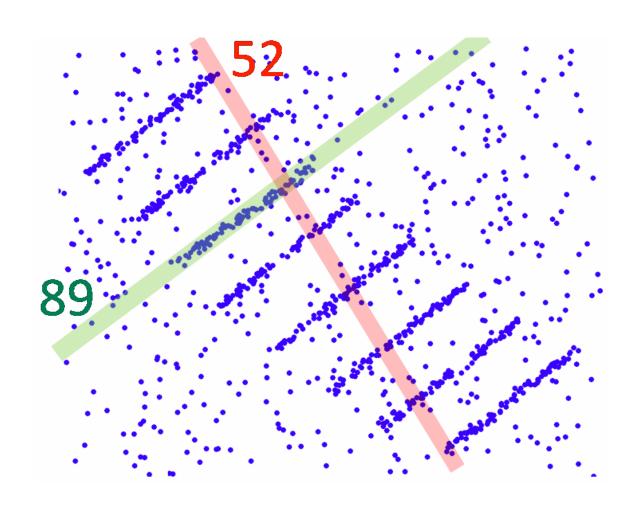


Why not RANSAC again?

remove inliers for line 1 and use RANSAC again (sequential RANSAC)

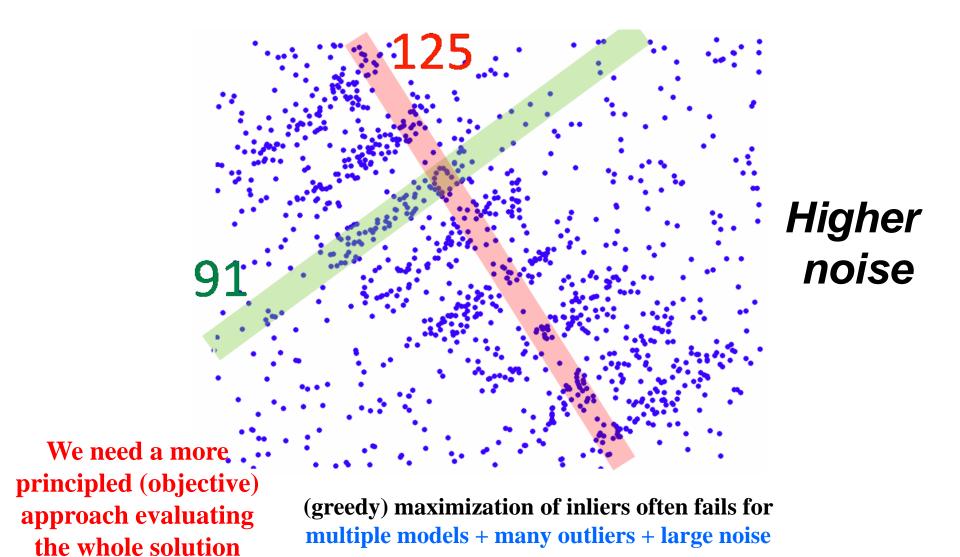


Multiple models and many outliers





Multiple models and many outliers



RANSAC as optimization (of certain objective)

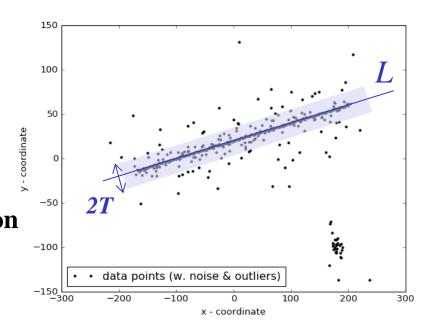


$$E(L) = \sum_{p} ||p - L||$$

- find optimal label L (line) minimizing loss function E(L)

RANSAC minimizes error (loss) function ||p-L|| counting outliers (equivalent to maximizing inliers)

treat ||p - L|| as an "operator" $|| \cdot - \cdot ||$ representing some error penalty function Loss(distance(p, L))



outliers inliers outliers
$$T \qquad dis$$

dist(p, L) - errors (distances between point p and line L)

RANSAC as optimization (of certain objective)

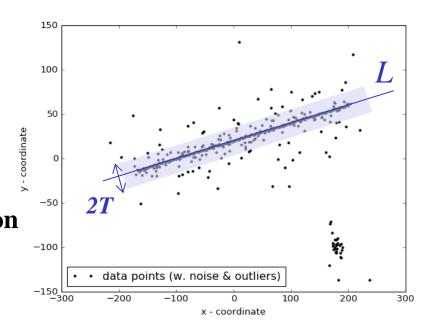


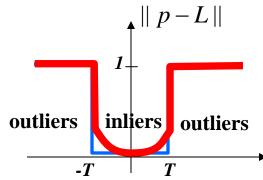
$$E(L) = \sum_{p} (||p - L||)$$

- find optimal label L (line) minimizing loss function E(L)

RANSAC minimizes error (loss) function ||p-L|| counting outliers (equivalent to maximizing inliers)

treat ||p - L|| as an "operator" $|| \cdot - \cdot ||$ representing some error penalty function Loss(distance(p, L))





The use of least squares among inliers can be interpreted as quadratic loss for errors < T

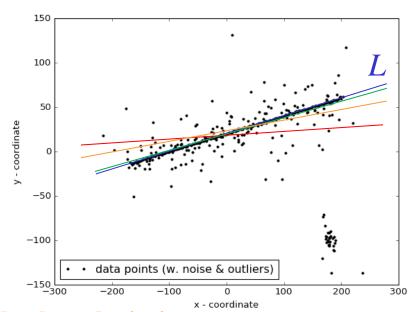
dist(p, L) - errors (distances between point p and line L)



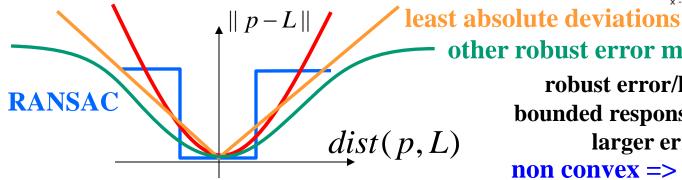
General error functions (overview)

$$E(L) = \sum_{p} ||p - L||$$

Can use different losses *i.e.* error functions ||p-L||



least squares



other robust error measure

robust error/loss functions have bounded response effectively ignoring larger errors (outliers)

non convex => harder to optimize

e.g. see iterative reweighted least squares (IRLS)



Error / loss function

distances to lines

$$E(\mathbf{L}) = \sum_{p} ||p - L_p||$$

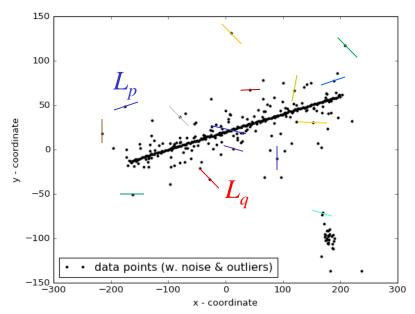
If multiple models

- assign different models (labels L_p) to every point p
 - find optimal labeling

$$\mathbf{L} = \{ L_1, L_2, \dots, L_n \}$$

$$\forall p, \ L_p \in \Lambda$$
set of used

models (lines)



too "complex" - too many lines (perfect line model for every point)

need a "simpler" solution, i.e. **few lines "explaining" all data**



Error / loss function

distances to lines

$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_{p}|| + \gamma \cdot |\Lambda|$$

Penalize the number of models (solution "complexity") $\gamma \cdot |\Lambda| \quad \text{our first example} \\ \text{of } \textit{regularization}$

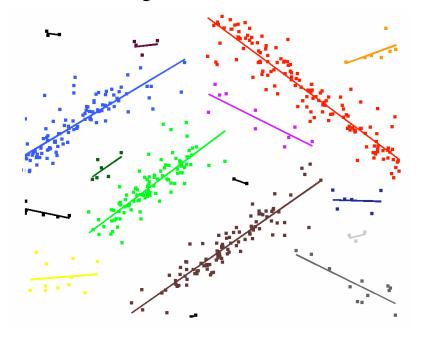
[Delong, Isack, et al. 2012]

If multiple models

- assign different models (labels L_p) to every point p
 - find optimal labeling

$$\mathbf{L} = \{ L_1, L_2, ..., L_n \}$$

$$\forall p, L_p \in \Lambda$$
set of used models (lines)



few lines "explaining" all data



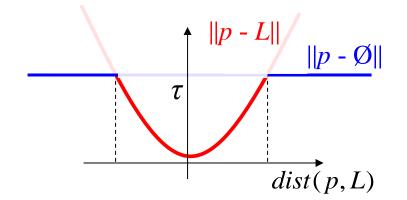
Error / loss function

distances to lines

$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_{p}|| + \gamma \cdot |\Lambda|$$

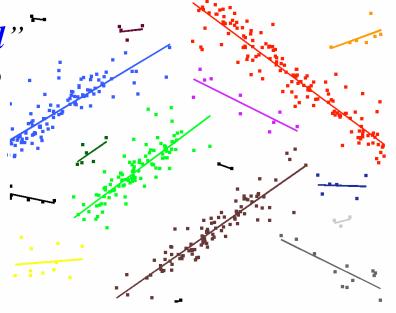
Extra trick: add to Λ one special model \emptyset representing "outlier model" with fixed cost $||p - \emptyset|| = \tau$ for any p

Then any point p prefers to be assigned "outlier model" \emptyset if its closest line L has $||p - L|| > \tau$



Penalize the number of models (solution "complexity") $\gamma \cdot |\Lambda| \quad \text{our first example} \\ \text{of } \textit{regularization}$

[Delong, Isack, et al. 2012]





Error / loss function

distances to lines

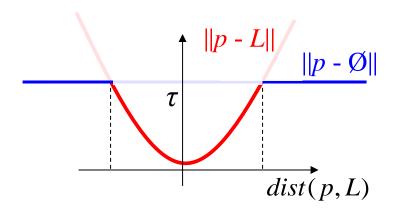
$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_{p}|| + \gamma \cdot |\Lambda|$$

Penalize the number of models (solution "complexity") $\gamma \cdot |\Lambda| \quad \text{our first example} \\ \text{of } \textit{regularization}$

[Delong, Isack, et al. 2012]

Extra trick: add to Λ one special model \emptyset representing "outlier model" with fixed cost $||p - \emptyset|| = \tau$ for any p

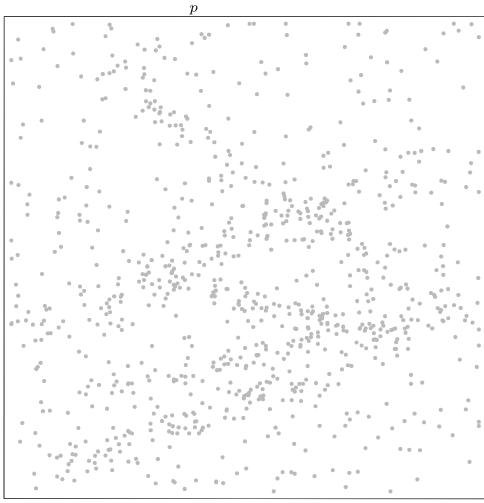
Then any point p prefers to be assigned "outlier model" \emptyset if its closest line L has $||p - L|| > \tau$



Also, weak models with only a few inliers $N < \frac{\gamma}{\tau}$ are not worth having. Its inliers should be assigned Ø. Why?



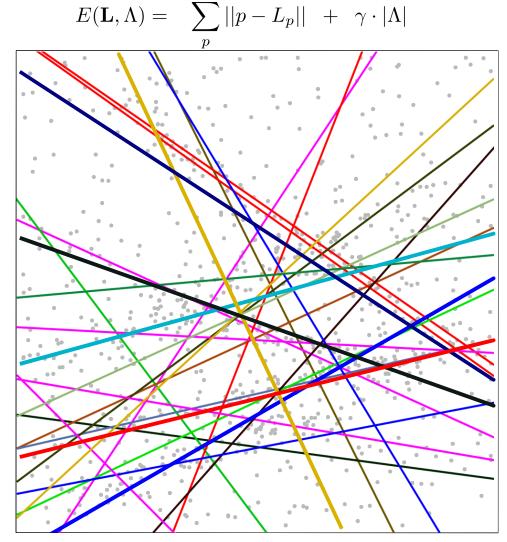
$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_p|| + \gamma \cdot |\Lambda|$$



data points



1. Initialization of Λ: randomly sampleK lines from points (some very large K)



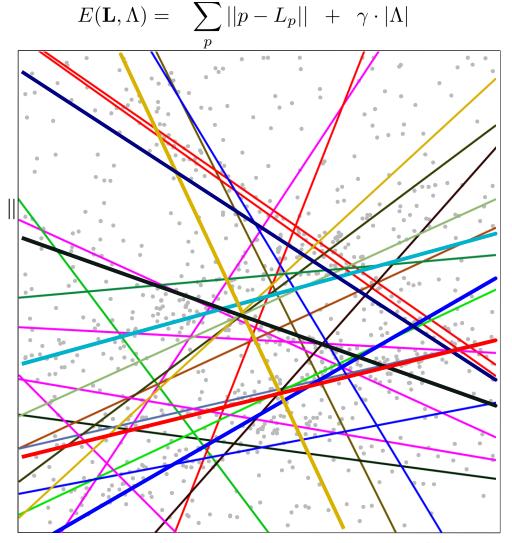
data points + randomly sampled lines



1. Initialization of Λ : randomly sample

K lines from points (some very large K)

2a. Assign each point to model in Λ with lowest $\| \|$



data points + randomly sampled lines

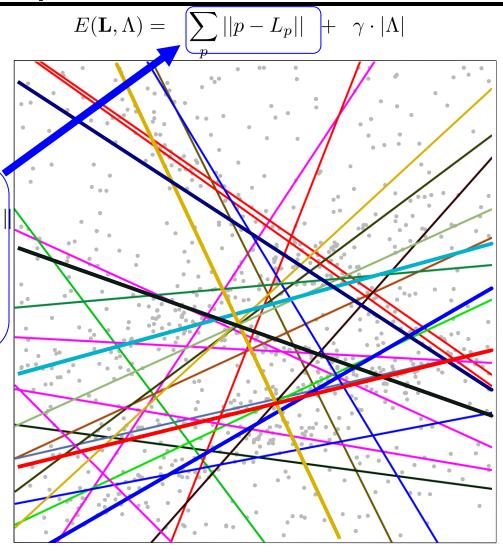


1. Initialization of Λ: randomly sampleK lines from points (some very large K)

2a. Assign each point to model in Λ with lowest $\parallel \parallel$ **2b.** Re-estimate lines parameters minimizing errors $\parallel \parallel$ among inliers (e.g. least squares)

addresses the first term of the objective assuming the number of used models is fixed:

$$|\Lambda| = const$$



data points + randomly sampled lines



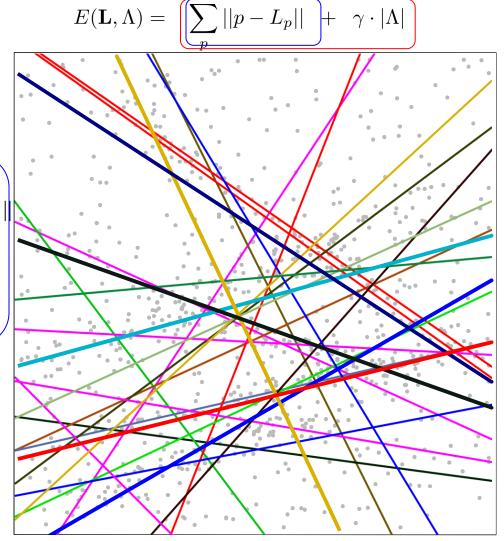
E relates to the objective for Uncapacitated Facility Location (UFL) problem Optimization problem

1. Initialization of Λ : randomly sample K lines from points (some very large K)

2a. Assign each point to model in Λ with lowest $\|\cdot\|$ **2b. Re-estimate** lines parameters minimizing

errors || || among inliers

(e.g. least squares)



data points + randomly sampled lines

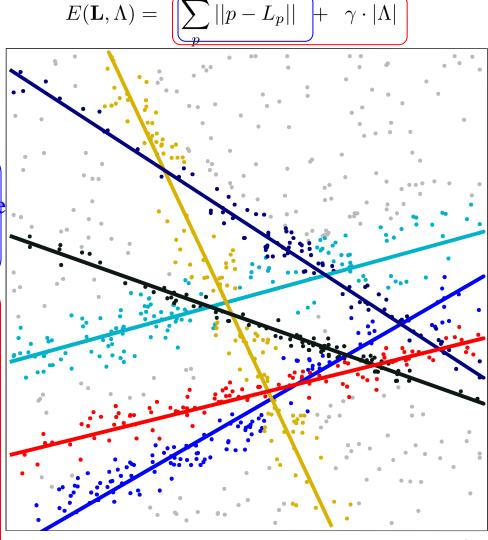


1

Optimization problem

- 1. Initialization of Λ: randomly sampleK lines from points(some very large K)
- 2. Assign points to the closest model, re-estimate line parameters using inliers (e.g. least squares)
- 3. (UFL heuristic) clean Λ by removing lines L that are not worth keeping:

$$\sum_{p:L_p=L} \min_{l \in \Lambda \setminus L} \|p - l\| < \sum_{p:L_p=L} \|p - L\| + \gamma$$



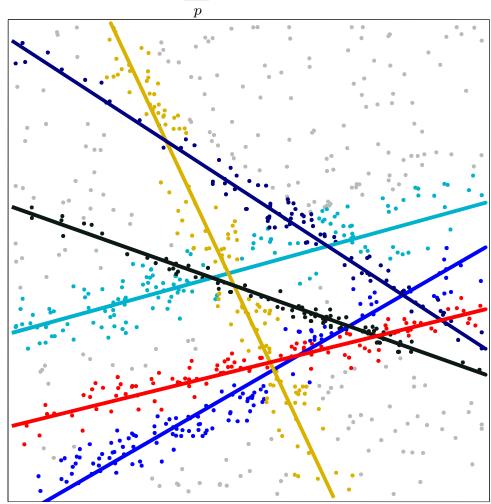
data points & assigned models (lines or Ø)

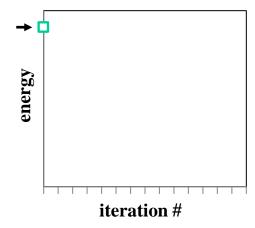
4. Iterate steps 2-3 until convergence



Optimization problem (algorithm illustration)

$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_p|| + \gamma \cdot |\Lambda|$$

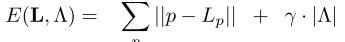


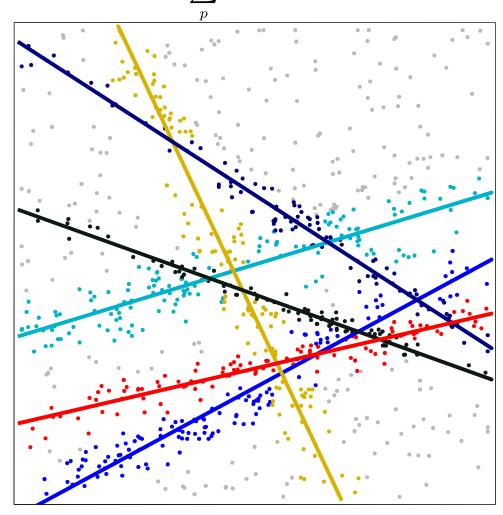


Beginning of iteration 2: points assigned to the closest model that survived iteration 1



Optimization problem (algorithm illustration)





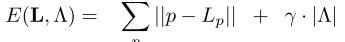
Re-estimating
lines in
↑
from their inliers
Remove

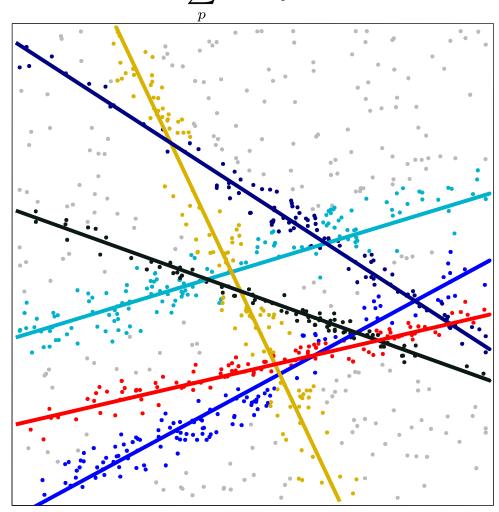
suboptimal lines, if any (UFL heuristic)

energy iteration #

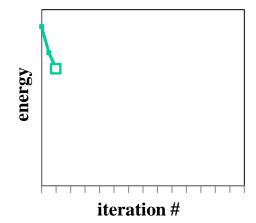
iteration 2: re-estimate model parameters







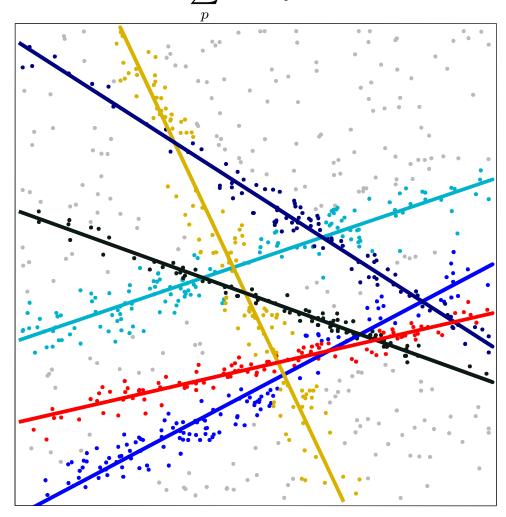
Re-assign points to closest lines



iteration 3: optimize points labeling L

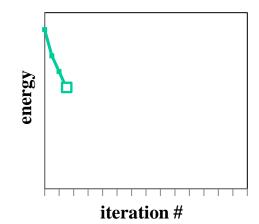


$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_p|| + \gamma \cdot |\Lambda|$$



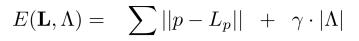
Re-estimating lines in from their inliers

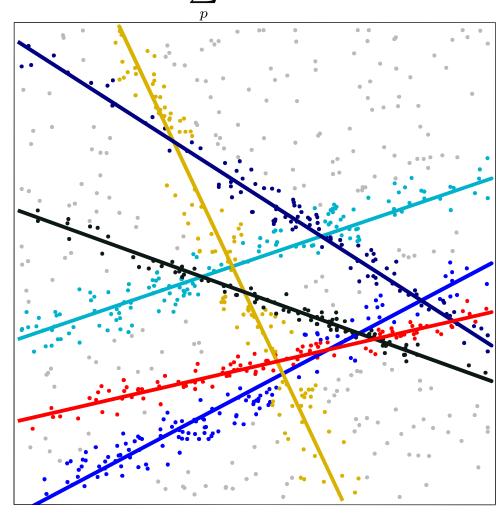
Remove suboptimal lines, if any (UFL heuristic)



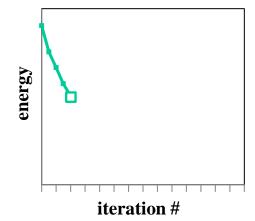
iteration 3: re-estimate model parameters







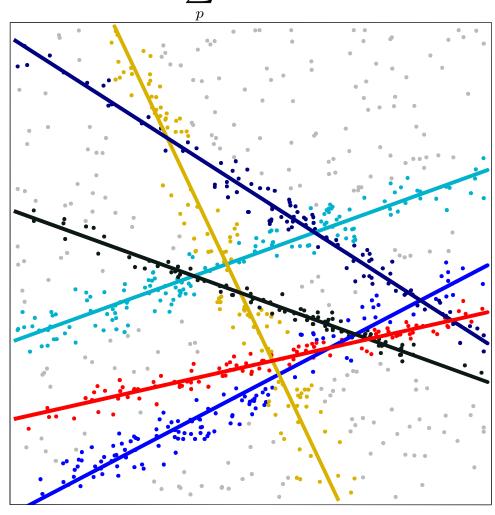
Re-assign points to closest lines



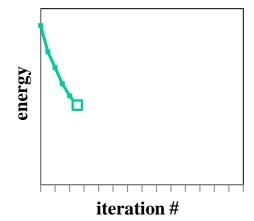
iteration 4: optimize points labeling L



$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_p|| + \gamma \cdot |\Lambda|$$



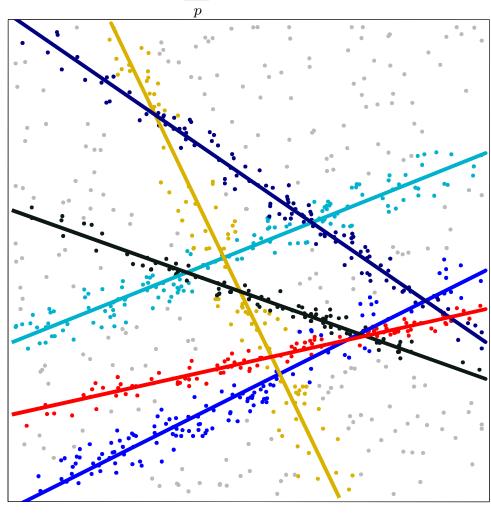
Re-estimating lines in from their inliers



iteration 4: re-estimate model parameters



$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_p|| + \gamma \cdot |\Lambda|$$

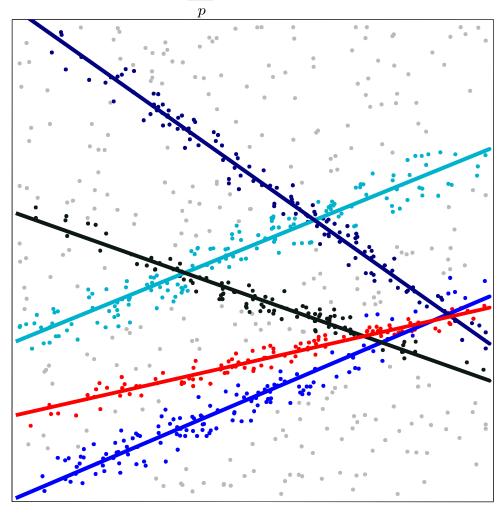


iteration #

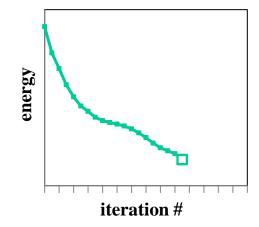
iteration 7...



$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_p|| + \gamma \cdot |\Lambda|$$



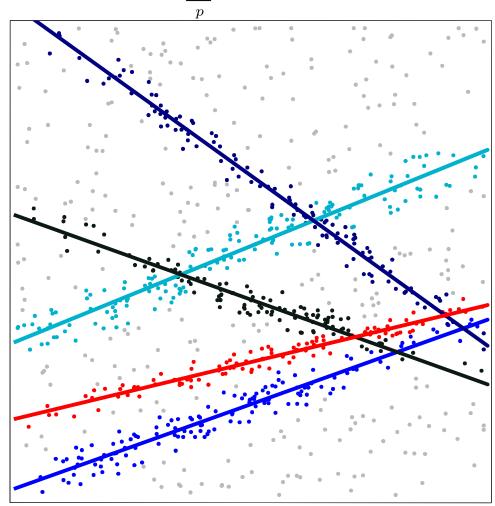
NOTE: yellow line became unnecessary

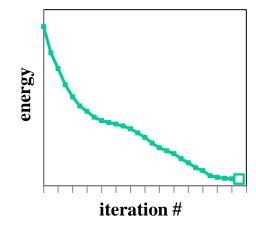


iteration 10...



$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_p|| + \gamma \cdot |\Lambda|$$

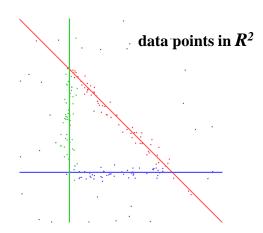




iteration 15... converged.

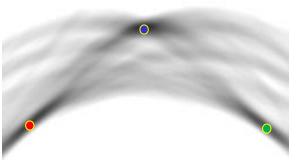
What's going on inside the space of labels (lines/models)







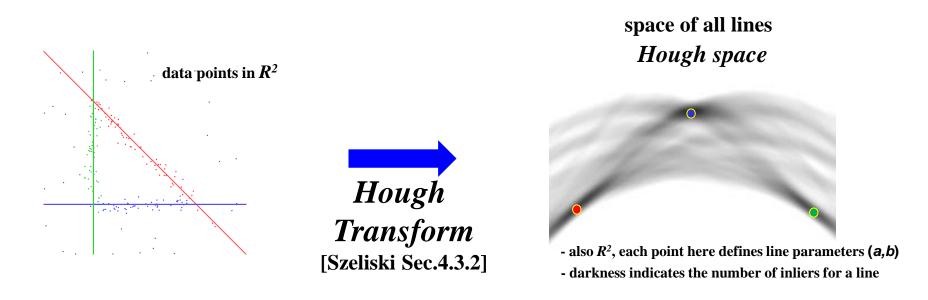
space of all lines Hough space



- also R^2 , each point here defines line parameters (a,b)
- darkness indicates the number of inliers for a line

What's going on inside the space of labels (lines/models)



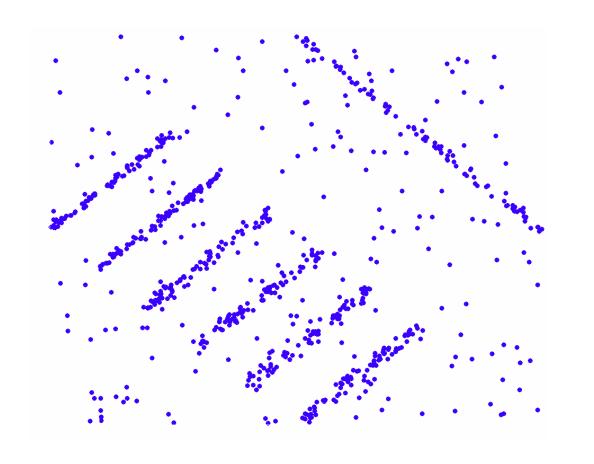


Q: why not just look for *modes* in the Hough space?

(strong local maxima)

Note: RANSAC searches the maxima by exploring a (large) sample of lines randomly sampled from the "line density" in the Hough space.

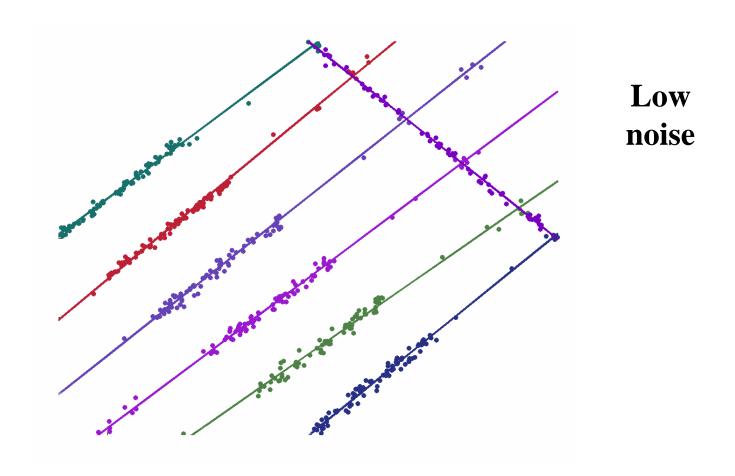




Low noise

original data points

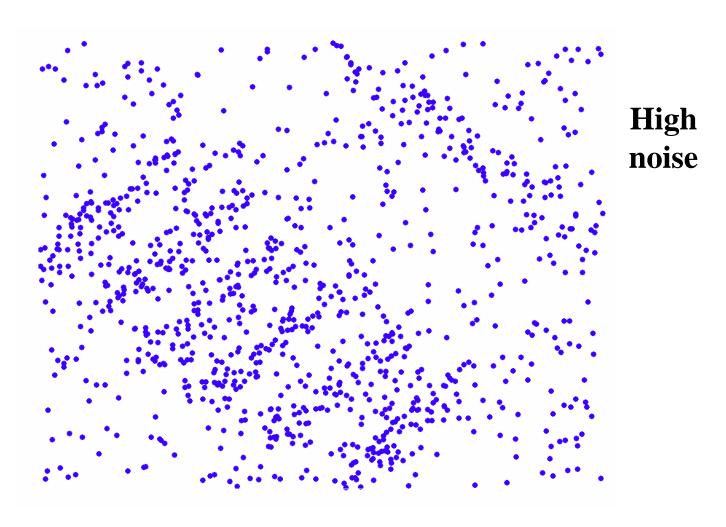




sequential RANSAC, modes in Hough space, UFL approach



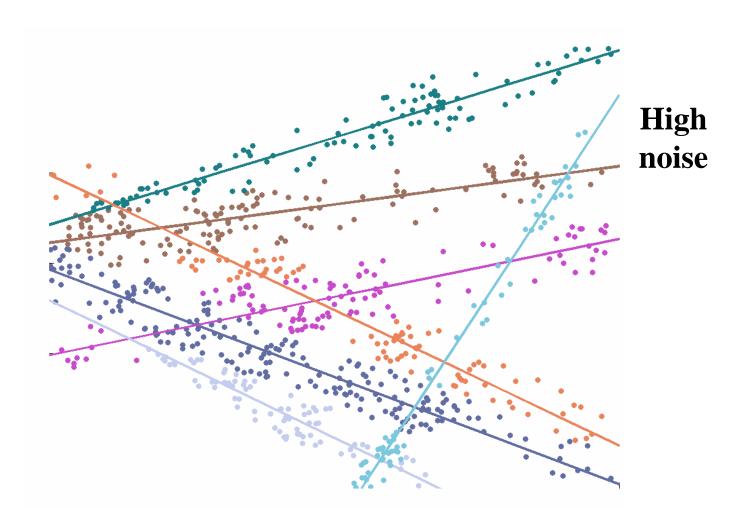




original data points



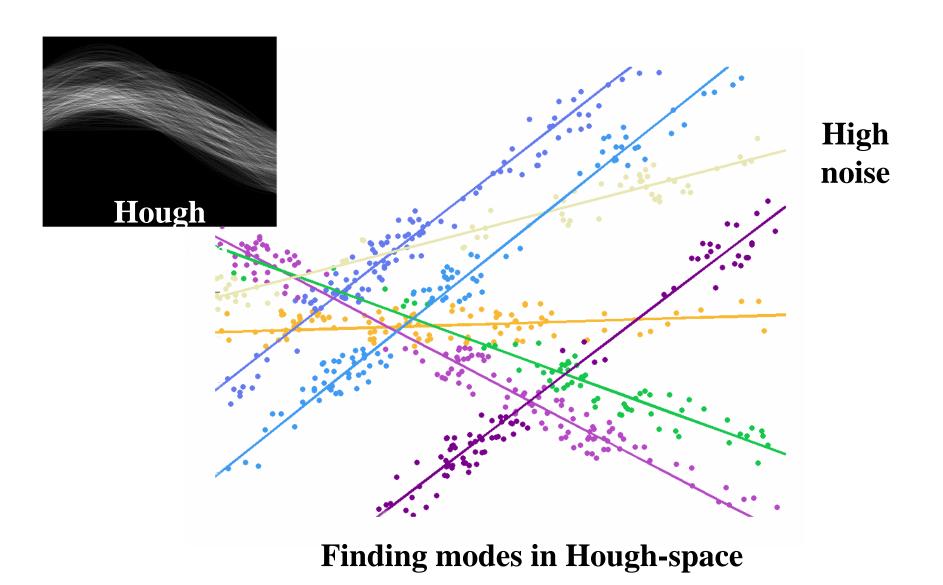




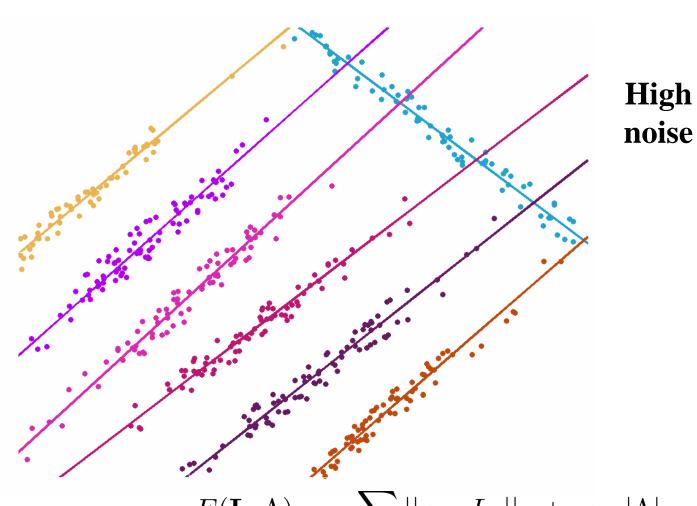
sequential RANSAC



Comparison



Comparison

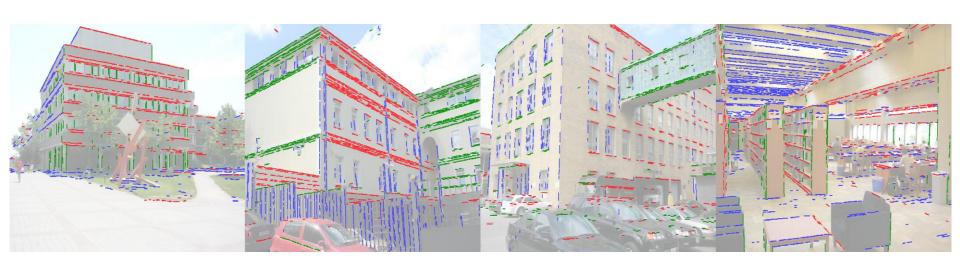


UFL-based approach [Delong et al. IJCV12]

$$E(\mathbf{L}, \Lambda) = \sum_{p} ||p - L_p|| + \gamma \cdot |\Lambda|$$



Line fitting on real image data



data points are "Canny edges"

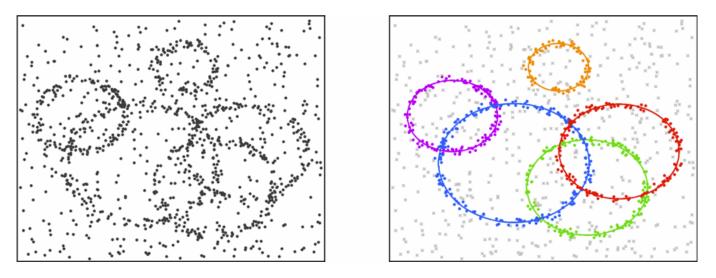
[Delong et al. NIPS 2012]

Note: color indicates clusters of lines with common vanishing point



Fitting other geometric models

$$||p-\theta|| := ((x_p - c_x)^2 - (y_p - c_y)^2 - r^2)^2$$
 for $\theta = \{c_c, c_y, r\}$



Model fitting for arbitrary geometric models θ

Need: 1) define an error measure w.r.t. model parameters $\|p - \theta\|$

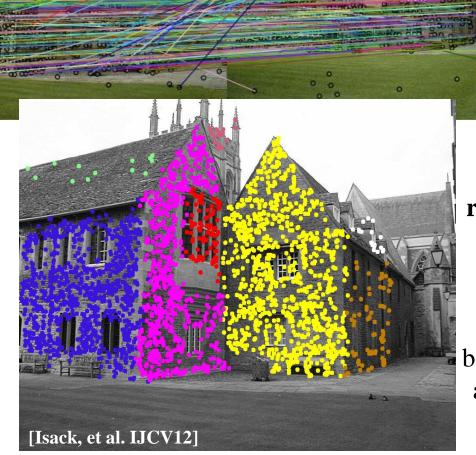
2) efficient method for minimizing the sum of errors among inliers w.r.t. model parameters θ

$$\min_{\theta} \sum_{p \in S} \| p - \theta \|$$



Fitting multiple homographies (e.g. planes)





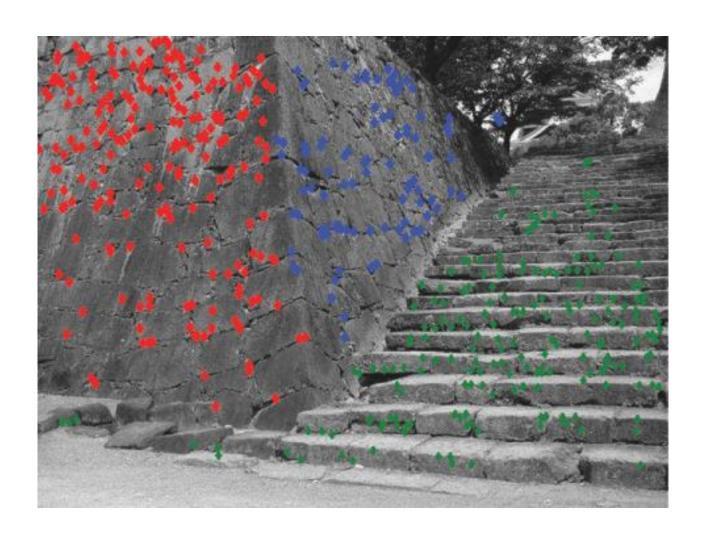
using
symmetric
re-projection errors

$$||p'-Hp||+||p-H^{-1}p'||$$

as an error measure between match (p, p')and homography H

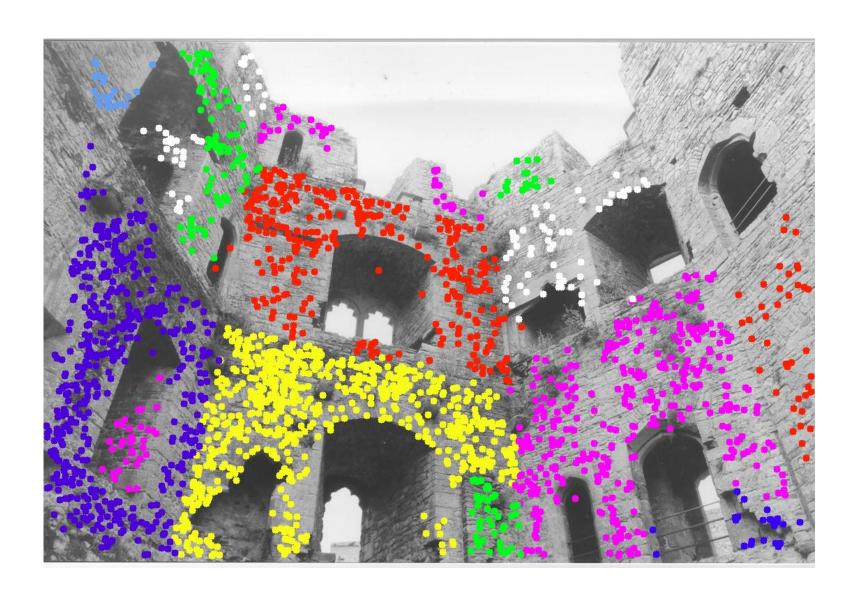


Fitting multiple homographies (e.g. planes)





Fitting multiple homographies (e.g. planes)



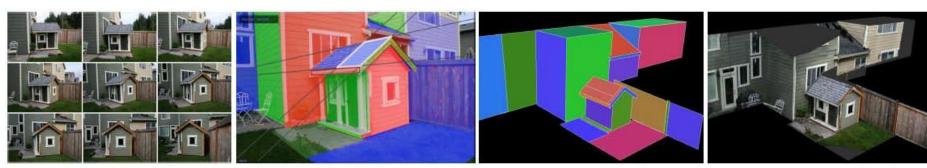


same scene from a different view point...



Note very small steps between each floor





Input Photographs 2D Sketching Interface Geometric Model Texture-mapped model

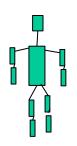
user can help with rough initialization instead of random sampling

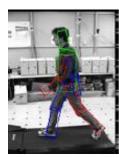
[Sinha et al. 2008]

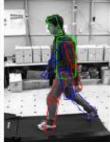


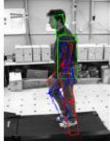
Fitting dependent models (parts)

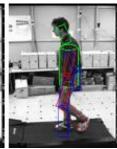
Pictorial structures [Felzenswalb & Huttenlocher, 2005]











Articulated model tracking

hand model fitting to 3D point cloud

















[Andrea Tagliasacchi et al., 2013]

■ Finding low dim. subspaces

e.g. in the space of point tracks



[Carl Olsson et al.,2017]



Fitting dependent models (parts)

Can use **physics** to define objectives (losses):

- parts interactions (e.g. spring-like)
- kinematics
- attraction/repulsion to image features

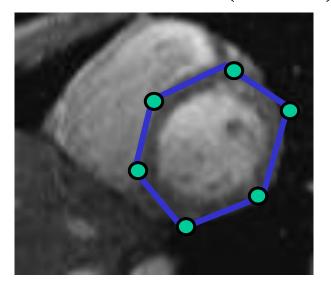
Articulated model tracking

hand model fitting to 3D point cloud



[Andrea Tagliasacchi et al., 2013]

Active contours (snakes)



[Kass Witkin Terzopoulos, 1988]



Geometric model fitting in vision

MODELS: lines, planes, homographies, affine transformations, projection matrices, fundamental/essential matrices, etc.

next topic

- single models (e.g. panorama stitching, camera projection matrix)
- multiple models (e.g. multi-plane reconstruction, multiple rigid motion)

FIRST STEP: detect some features (corners, LOGS, etc) and compute their descriptors (SIFT, MOPS, etc.)

SECOND STEP: match or track

THIRD STEP: fit models

(minimization or errors/losses)