## Geometric Model Fitting


with slides stolen from
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## Geometric Model Fitting

- Feature matching $\left(\mathbf{p}_{i}, \mathbf{p}_{i}^{\prime}\right)$
- 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 <br> 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

$8 \times 8$ oriented patch

- Sampled at $5 \times$ scale

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


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

## Flashbacks: MOPS descriptor

$8 \times 8$ oriented patch

- Sampled at $5 \times$ scale

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


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

## Feature matching



## Feature matching

## 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 $\operatorname{SSD}$ (patch1, patch2) $<T$ (threshold)


## Feature matching

SSD(patch1, patch2) < T

How to set threshold T?

> no threshold T is good for separating correct and incorrect matches

## Feature matching

A better way [Lowe, 1999]:

- SSD of the closest match (SSD1)
- SSD of the second-closest 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 $(\mathrm{imL})$, threshold_rel $=0.0005$, min_distance $=5$ )
keypointsR $=$ corner_peaks $($ corner_harris $(i m R)$, 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]
find the closest match $p$, for any feature $p$
descriptorsR $=$ extractor.descriptors

## 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 $\left(\mathbf{p}_{i}, \mathbf{p}_{i}^{\prime}\right)$ is more than 4 Answer: model fitting via "least squares" (later, slide 21)

Issue 2: too many outliers or wrong matches $\left(\mathbf{p}_{i}, \mathbf{p}_{i}^{\prime}\right)$
Answer: robust model fitting via RANSAC (later, slide 35)

## Recall: Homography from 4 points



## Recall: Homography from 4 points

Consider one match (point-correspondence) $p=(x, y) \rightarrow p^{\prime}=\left(x^{\prime}, y^{\prime}\right)$

$$
\mathbf{p}^{\prime}=\mathbf{H p} \quad\left[\begin{array}{c}
w x^{\prime} \\
w y^{\prime} \\
w
\end{array}\right]=\left[\begin{array}{lll}
a & b & c \\
d & e & f \\
g & h & i
\end{array}\right]\left[\begin{array}{c}
x \\
y \\
1
\end{array}\right]
$$

After eliminating $w=g x+h y+i$ :

$$
\Rightarrow \quad \begin{aligned}
& a x+b y+c-g x x^{\prime}-h y x^{\prime}-i x^{\prime}=0 \\
& d x+e y+f-g x y^{\prime}-h y y^{\prime}-i y^{\prime}=0
\end{aligned}
$$

Two equations linear w.r.t unknown coefficients of matrix $H$ and quadratic w.r.t. known point coordinates ( $x, y, x^{\prime}, y^{\prime}$ )

## Recall: Homography from 4 points

Consider 4 point-correspondences $p_{i}=\left(x_{i}, y_{i}\right) \rightarrow p_{i}^{\prime}=\left(x_{i}^{\prime}, y_{i}^{\prime}\right)$

$$
\left.\left.\left.\begin{array}{r}
\mathbf{p}_{i}^{\prime}=\mathbf{H p}_{i} \\
w_{i} y_{i}^{\prime} \\
w_{i}
\end{array}\right]=\left[\begin{array}{ccc}
w_{i} x_{i}^{\prime} \\
d & e & f \\
g & h & i
\end{array}\right]\left[\begin{array}{ccc}
a & b & c \\
y_{i} \\
1
\end{array}\right] \quad \begin{array}{c}
x_{i} \\
\text { for i=1,2,3,4 }
\end{array}\right] \begin{array}{c}
\text { Special case of } \\
\text { DLT method } \\
\text { (see p.89 } \\
\text { in Hartley and } \\
\text { Zisserman) }
\end{array}\right]
$$

Can be written as matrix multiplication $\quad \mathbf{A}_{i} \cdot \mathbf{h}=\mathbf{0} \quad$ for $\mathbf{i}=\mathbf{1 , 2 , 3 , 4}$
where $\mathbf{h}=[a b c d e f g h i]^{T}$ is a vector of unknown coefficients in $\mathbf{H}$ and $\mathbf{A}_{i}$ is a $2 \times 9$ matrix based on known point coordinates $x_{i}, y_{i}, x_{i}^{\prime}, y_{i}^{\prime}$

## Recall: Homography from 4 points

Consider 4 point-correspondences $p_{i}=\left(x_{i} y_{i}\right) \rightarrow p_{i}^{\prime}=\left(x_{i}^{\prime}, y_{i}^{\prime}\right)$

$$
\mathbf{p}_{i}^{\prime}=\mathbf{H} \mathbf{p}_{i} \quad \Rightarrow \quad \underset{2 \times 99 \times 1 \quad 2 \times 1}{\mathbf{A}_{i} \cdot \mathbf{h}=\mathbf{0}} \quad \text { for } \mathrm{i}=1,2,3,4
$$

## Recall: Homography from 4 points

Consider 4 point-correspondences $p_{i}=\left(x_{i} y_{i}\right) \rightarrow p_{i}^{\prime}=\left(x_{i}^{\prime}, y_{i}^{\prime}\right)$

$$
\begin{aligned}
& \mathbf{p}_{i}^{\prime}=\mathbf{H} \mathbf{p}_{i} \\
& \text { for } \mathrm{i}=1,2,3,4
\end{aligned}
$$

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 7,
To find one specific solution $h$, for now fix one element, e.g. $i=1$ this may not work more generally, should


## Recall: Homography from 4 points

Consider 4 point correspondences $p_{i}=\left(x_{i} y_{i}\right) \rightarrow p_{i}^{\prime}=\left(x_{i,}^{\prime}, y_{i}^{\prime}\right)$

$$
\begin{aligned}
& \mathbf{p}_{i}^{\prime}=\mathbf{H} \mathbf{p}_{i} \\
& \text { for } i=1,2,3,4
\end{aligned} \quad \Rightarrow \quad{\underset{8 \times 8}{\mathbf{A}_{1: 8}} \cdot \mathbf{h}_{1: 8}=-\mathbf{A}_{9 \times 1}}_{8 \times 1}
$$

## More than 4 points

Consider 4 point correspondences $p_{i}=\left(x_{i}, y_{i}\right) \rightarrow p_{i}^{\prime}=\left(x_{i,}^{\prime} y_{i}^{\prime}\right)$

$$
\begin{gathered}
\mathbf{p}_{i}^{\prime}=\mathbf{H} \mathbf{p}_{i} \\
\text { for } \mathrm{i}=1,2,3,4
\end{gathered} \quad \Rightarrow \quad \begin{array}{|c}
\mathbf{A}_{1: 8} \cdot \mathbf{h}_{1: 8} \\
8 \times 8
\end{array}=-\mathbf{A}_{9}
$$

## Questions:

What if 4 points correspondences are known with error?

Are there any benefits from knowing more point correspondences?

First, consider a simpler model fitting problem...

## Simpler example: line fitting

Assume a set of data points $\left(X_{1}, X_{1}^{\prime}\right),\left(X_{2}, X_{2}^{\prime}\right),\left(X_{3}, X_{3}^{\prime}\right), \ldots$
(e.g. person's height vs. weight)

We want to fit a model (e.g. a line) to predict $X^{\prime}$ from $X$

$$
a \cdot X+b=X^{\prime}
$$

How many pairs ( $X_{i}, X_{i}^{\prime}$ ) do we need to find $a$ and $b$ ?

$$
\begin{gathered}
X_{1} a+b=X_{1}^{\prime} \\
X_{2} a+b=X_{2}^{\prime} \\
{\left[\begin{array}{ll}
X_{1} & 1 \\
X_{2} & 1
\end{array}\right]\left[\begin{array}{l}
a \\
b
\end{array}\right]=\left[\begin{array}{l}
X_{1}^{\prime} \\
X_{2}^{\prime}
\end{array}\right]} \\
\boldsymbol{A} \cdot \boldsymbol{x}=\boldsymbol{B}
\end{gathered} \boldsymbol{x}^{\boldsymbol{x}=\boldsymbol{A}^{-1} \cdot \boldsymbol{B}} .
$$



## Simpler example: line fitting

Assume a set of data points $\left(X_{1}, X_{1}^{\prime}\right),\left(X_{2}, X_{2}^{\prime}\right),\left(X_{3}, X_{3}^{\prime}\right), \ldots$
(e.g. person's height vs. weight)

We want to fit a model (e.g. a line) to predict $X^{\prime}$ from $X$

$$
a \cdot X+b=X^{\prime}
$$

What if the data points ( $X_{i}, X_{i}^{\prime}$ ) are noisy?

$$
\begin{aligned}
& {\left[\begin{array}{cc}
X_{1} & 1 \\
X_{2} & 1 \\
X_{3} & 1 \\
\ldots & \ldots
\end{array}\right]\left[\begin{array}{l}
a \\
b
\end{array}\right]=\left[\begin{array}{c}
X_{1}^{\prime} \\
X_{2}^{\prime} \\
X_{3}^{\prime} \\
\ldots
\end{array}\right]} \\
& \min _{x}\|A x-B\|^{2} \\
& \boldsymbol{x}=\boldsymbol{A}^{-1} \cdot \boldsymbol{B}
\end{aligned}
$$

$$
\begin{array}{ll} 
& \text { this problem is also known as } \\
\chi^{\prime} & \begin{array}{l}
\text { "linear regression" problem }
\end{array}
\end{array}
$$

$$
a \cdot X+b=X^{\prime}
$$

$$
\min _{x}\|A x-B\|^{2} \quad \text { (least-squares) }
$$

$$
\text { where } \quad A^{-1} \equiv V \cdot W^{-1} \cdot U^{T} \quad \text { is a pseudo-inverse }
$$

$$
\begin{aligned}
& \text { where } A \equiv V \cdot W \cdot \text { is a pseudo-inv } \\
& \text { based on SVD decomposition } A=U \cdot W \cdot V^{T}
\end{aligned}
$$

(in python, one can use $s v d$ function in library numpy.linalg)

## SVD: rough idea

$$
\underset{\mathrm{M} \geq \mathrm{N}:}{A}=\underbrace{\text { embed }}_{\mathrm{MxN}} \cdot V_{\mathrm{NxN}}^{\text {scale }} \cdot V_{\mathrm{NxN}}^{\text {route }} T
$$

where $U$ and $V$ are matrices with orthonormal columns and $W$ is diagonal with elements $\boldsymbol{w}_{i} \geq 0$ (see "Numerical Recipes in C", edition 2, Sec. 2.6)


 3x2


Q: where are the points from $R^{\mathbf{2}}$ mapped to? A: point B: line C: plane D: whole $\mathrm{R}^{3}$


How does SVD help to solve least-squares?

$$
\min _{x}\|A x-B\|^{2}
$$

projection of $B$ onto range of $\boldsymbol{A}$


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

Equivalent (fast to compute) expression

$$
\left.\underset{\mathrm{Nx} 1}{x}=\underset{\mathrm{NxN}}{\left(A^{T}\right.} \boldsymbol{A}\right)^{-1} \cdot \underset{\mathrm{NxM}}{A^{T}} \cdot \underset{\mathrm{Mx1}}{B}
$$

Indeed: $\left.\boldsymbol{A}^{\boldsymbol{T}} \boldsymbol{A}=V W U^{2} \not\right)^{2} W V^{T}=\boldsymbol{V} \boldsymbol{W}^{2} \boldsymbol{V}^{\boldsymbol{T}}$ so $\left(\boldsymbol{A}^{\boldsymbol{T}} \boldsymbol{A}\right)^{-1} \cdot \boldsymbol{A}^{\boldsymbol{T}}=V W^{-2}, ~ N W U^{T}=\boldsymbol{V} \boldsymbol{W}^{-1} \boldsymbol{U}^{\boldsymbol{T}}$ If $\mathrm{M} \gg \mathrm{N}$ computing inverse of positive semi-definite NuN matrix $A^{T} A$ can be faster than SVD of MuN matrix A

## Least squares line fitting

Data generated as $X_{i}^{\prime}=a X_{i}+b+\delta X_{i}$ for $a=0.2, b=20$ and Normal noise $\delta X_{i}$


## Least squares fail in presence of outliers

Data generated as $X_{i}^{\prime}=a X_{i}+b+\delta X_{i}$ for $a=0.2, b=20$ and Normal noise $\delta X_{i}$


For cases with outliers we need a more robust method (e.g. RANSAC, coming soon)

## Model fitting robust to outliers

We need a method that can separate inliers from outliers

## RANSAC

random sampling
consensus
[Fischler and Bolles, 1981]

## RANSAC (for line fitting example)



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

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2. count inliers $p$ for threshold $T$

$$
\|p-l\| \leq T
$$

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$$
\|p-l\| \leq T
$$

3. repeat

## RANSAC (for line fitting example)



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2. count inliers $p$ for threshold $T$

$$
\|p-l\| \leq T
$$

3. repeat $\mathbf{N}$ times and select model with most inliers

## RANSAC (for line fitting example)



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

$$
\|p-l\| \leq T
$$

3. repeat $\mathbf{N}$ 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 are both inliers? [Fischler and Bolles, 1981]

## RANSAC (for line fitting example)



$$
\boldsymbol{X}_{i} \quad \text { (similar math as in statistical analysist of RANSAC success rate) }
$$

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

## So, how do we find multiple models?



## So, how do we find multiple models?



## Homography from $N \geq 4$ points

Consider N point correspondences $p_{i}=\left(x_{i}, y_{i}\right) \rightarrow p_{i}^{\prime}=\left(x_{i}^{\prime}, y_{i}^{\prime}\right)$

$$
\begin{aligned}
& \mathbf{p}_{i}^{\prime}=\mathbf{H} \mathbf{p}_{i} \\
& \text { for } i=1, \ldots, \mathrm{~N}
\end{aligned}
$$



$$
\underset{2 N \mathrm{~N} \cdot}{\mathbf{A} \cdot \mathbf{h}=\mathbf{0}=\mathbf{0}}
$$

over-constrained system

Approach 1: add constraint $\boldsymbol{i}=\mathbf{1}$. So, there are only 8 unknowns.
Set up a system of linear equations for vector of unknowns $\boldsymbol{h}_{1: 8}=[\mathrm{a}, \mathrm{b}, \mathrm{c}, \mathrm{d}, \mathrm{e}, \mathrm{f}, \mathrm{g}, \mathrm{h}]^{\mathrm{T}}$

$$
\underset{\substack{2 \times 88}}{\mathbf{A}_{1: 8}} \cdot \underset{8 \mathrm{x}}{\mathbf{h}} \underset{1: 8}{ }=\mathbf{B}=-\underset{2 \mathrm{x} 1}{\mathbf{A}_{9}}
$$

solve

$$
\min _{h_{t s}}\left\|A_{1: 8} h_{1: 8}-B\right\|^{2}
$$

(least-squares)
compute inverse for $\mathbf{A}_{1: 8}^{T} \mathbf{A}_{1: 8}$ as in line fitting, then $\boldsymbol{h}_{\mathbf{1}: \mathbf{8}}=\left(\mathbf{A}_{\mathbf{1}: 8}{ }^{T} \cdot \mathbf{A}_{\mathbf{1}: 8}\right)^{-1} \cdot \mathbf{A}_{\mathbf{1}: 8}{ }^{T} \cdot\left(-\mathbf{A}_{9}\right)$

## Homography from $N \geq 4$ points

Consider N point correspondences $p_{i}=\left(x_{i}, y_{i}\right) \rightarrow p_{i}^{\prime}=\left(x_{i}^{\prime}, y_{i}^{\prime}\right)$

$$
\begin{aligned}
& \mathbf{p}_{i}^{\prime}=\mathbf{H} \mathbf{p}_{i} \\
& \text { for } i=1, \ldots, \mathrm{~N}
\end{aligned}
$$



$$
\underbrace{}_{2 N \times 9} \mathbf{A} \cdot \mathbf{h}=\mathbf{0}
$$

over-constrained system

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


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 $\quad W=\left[\begin{array}{cc}w_{1} & 0 \\ 0 & w_{2}\end{array}\right]$

solve: $\min _{x:\|x\|=1}\|W \cdot x\|$


Solution: $\quad x=(1,0)$ if $w_{1}<w_{2}$

$$
x=(0,1) \text { if } w_{2}<w_{1}
$$

$\Leftarrow$ equivalently, solve $\min _{u \in \text { Elips }}\|u\|$

## General case: use SVD (rough idea)


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

## General case: use SVD (rough idea)


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

$\uparrow \mathfrak{R}^{3}$


How does SVD help to solve $\min _{x:||x||=1}\|\mathbf{A} \cdot x\|$ ?

$$
A \cdot V_{i}=U W V^{T} \cdot V_{i}=U W \cdot e_{i}=w_{i} \cdot\left(U \cdot e_{i}\right)=w_{i} \cdot U_{i} \quad \Rightarrow\left\|A \cdot V_{i}\right\|=w_{i} \begin{aligned}
& \frac{\begin{array}{c}
\text { vector } x=V_{i} \\
\text { the least } w_{i} \\
\text { corresponding to } \\
\text { solves the problem }
\end{array}}{\text { then }} \text {. }
\end{aligned}
$$

Check that $V_{i}$ and $\left(w_{i}\right)^{2}$ are eigen vectors/values for matrix $A^{T} A$ (note: ellipsoid $x^{T} \cdot\left(A^{T} A\right) \cdot x=1$ maps onto circle $u^{T} \cdot u=1$ )
$A^{T} A \cdot V_{i}=V W^{2} V^{T} \cdot V_{i}=V W^{2} \cdot e_{i}=w_{i}^{2} \cdot V_{i} \quad \Rightarrow \quad$ can use eigen decomposition of $\boldsymbol{A}^{\boldsymbol{T}} \boldsymbol{A}$ instead of SVD of $\boldsymbol{A}$.

## Least squares fail in presence of outliers



Least squares work
if using "inliers" only (detecting these? - soon )

imR projected from inliers only


Least squares work if using "inliers" only (detecting these? - soon )

larger errors in the area with no matches
 we minimized

$a x+b y+c-g x x^{\prime}-h y x^{\prime}-i x^{\prime}$<br>$\approx 0$<br>$d x+e y+f-g x y^{\prime}-h y y^{\prime}-i y^{\prime}$<br>$\approx 0$

"geometric errors"
observable in the image

$$
\begin{aligned}
& x^{\prime}-\frac{a x+b y+c}{g x+h y+i} \approx 0 \\
& y^{\prime}-\frac{d x+e y+f}{g x+h y+i} \approx 0
\end{aligned}
$$

(harder to minimize)
"errors"
nong inliers $\left\|p^{\prime}-H p\right\| \quad$ Did we actually minimize these errors?

Least squares work if using "inliers" only (detecting these? - soon )


Question: how to remove outliers automatically?

## RANSAC for robust homography fitting


only two differences:

1. need to randomly sample four pairs $\left(p, p^{\prime}\right)$
the minimum number of matches to estimate a homography H
2. pair $\left(p, p^{\prime}\right)$ counts as an inlier for a given homography $\mathbf{H}$ if

$$
\left\|p^{\prime}-H p\right\| \leq T
$$

## RANSAC for robust homography fitting



Homography for corrupted four matches is likely to have only a few inliers ( $p, p^{\prime}$ )


$$
\left\|p^{\prime}-H p\right\| \leq T
$$

(randomly sampled)

## RANSAC for robust homography fitting



Homography for good four matches has 21 inliers ( $p, p^{\prime}$ )

$$
\left\|p^{\prime}-H p\right\| \leq T
$$

(randomly sampled)

## RANSAC for robust homography fitting



Inliers for the randomly sampled homography with the largest inlier set
imL in Reference frame (LtoRef)

imR projected from RANSAC inliers


## RANSAC for robust homography fitting


matched inliers
$\left\|p^{\prime}-H p\right\|$
The final automatic panorama result

## RANSAC for robust model fitting

RANSAC loop:
In general (for other models):
always sample the smallest number
of points/matches needed to estimate a model

1. Select four feature pairs (at random)
2. Compute homography H (exact)
3. Count inliers ( $p, p^{\prime}$ ) where $\left\|p^{\prime}-H p\right\| \leq T$
e.g. geometric errors
4. Iterate N times (steps $1-3$ ). Keep the largest set of inliers.
5. Re-compute least-squares H estimate on all of the inliers

## Other examples of geometric model fitting



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?

## Other examples of geometric model fitting



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?

## Other examples of geometric model fitting

1. Allow to remove bad matches (outliers)
2. Correspondences can be estimated with subpixel precision $\left(\boldsymbol{p}, \boldsymbol{p}^{\prime}\right) \rightarrow(\boldsymbol{p}, \boldsymbol{H} \boldsymbol{p})$
matched features ( $\mathrm{p}, \mathrm{p}$ '), as earlier


NOTE:
good matches can be used for reconstructing 3D points if camera positions \& orientations are known:
$\left(p, p^{\prime}\right) \rightarrow X_{p} \in R^{3}$
(Triangulation, see next topic)
3. Inliers allow to segment planar regions
4. Segmentation allows to extend correspondence from inliers to any point $\boldsymbol{p}$ inside segment: $(\boldsymbol{p}, \boldsymbol{H} \boldsymbol{p})$
5. Piece-wise planar 3D scene reconstruction

## So, how do we find multiple models?



## Fitting other geometric models

$$
\|p-\theta\|:=\left(\left(x_{p}-c_{x}\right)^{2}-\left(y_{p}-c_{y}\right)^{2}-r^{2}\right)^{2} \quad \text { for } \quad \theta=\left\{c_{c}, c_{y}, r\right\}
$$



Model fitting for arbitrary geometric models $\boldsymbol{\theta}$

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 $\boldsymbol{\theta}$

$$
\min _{\theta} \sum_{p \in S_{\theta}}\|p-\theta\|
$$

## Fitting multiple homographies (e.g. planes)



Fitting multiple homographies (e.g. planes)


## Fitting multiple homographies (e.g. planes)



## same scene from a different view point...



## 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)

