## Computer Vision

Class 05

Raquel Frizera Vassallo

### 2D Homography

Parameter Estimation

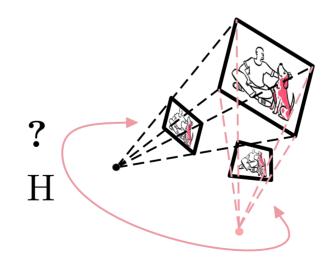
### Summary

- Homography
- DLT
- Normalized DLT
- Gold Standard Algorithm
- RANSAC
- Automatic computation of H using RANSAC



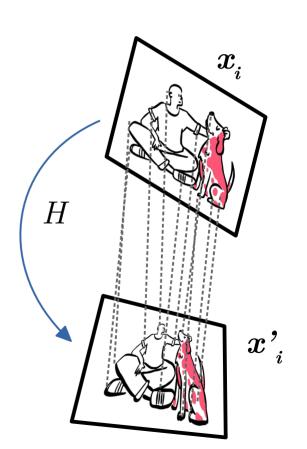
# **DLT**Direct Linear

Transformation



### Parameter Estimation

- 2D homography
- Given a set of  $(x_i, x_i')$ , compute  $H(x_i'=Hx_i)$
- Estimation will be based on the Direct Linear Transformation Algorithm (DLT)



### Number of measurements required

• At least as many independent equations as degrees of freedom required

$$x' = Hx$$

$$\lambda \begin{bmatrix} x' \\ y' \\ 1 \end{bmatrix} = \begin{bmatrix} h_{11} & h_{12} & h_{13} \\ h_{21} & h_{22} & h_{23} \\ h_{31} & h_{32} & h_{33} \end{bmatrix} \begin{bmatrix} x \\ y \\ 1 \end{bmatrix}$$

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- 8 degrees of freedom

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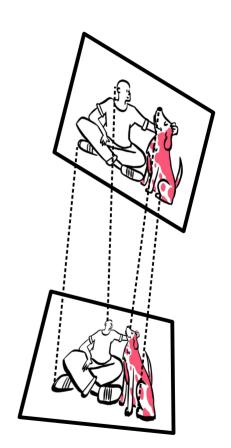
- 2 independent equations per point
- 8 degrees of freedom

We need 4 or more matchings

$$N \times 2 \ge 8 \rightarrow N \ge 4$$

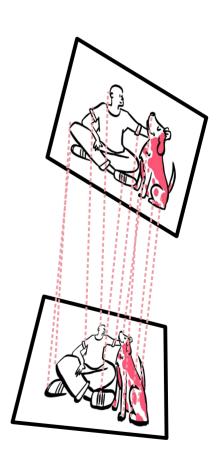
### Approximate solutions

- Minimal solution:
  - 4 points yield an exact solution for H.



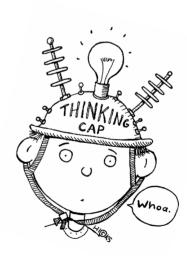
### Approximate solutions

- Minimal solution:
  - 4 points yield an exact solution for *H*.
- More points
  - No exact solution, because measurements are inexact ("noise").
  - Search for "best" according to some cost function.

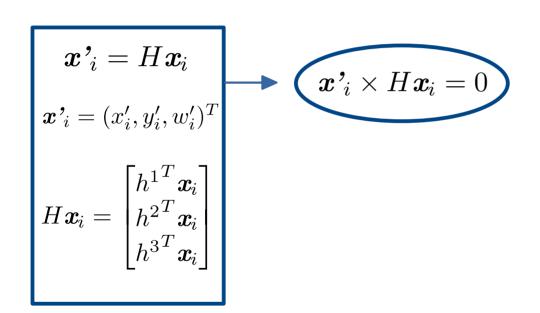


### Gold Standard algorithm

- Cost function that is optimal for some assumptions.
- Computational algorithm that minimizes it is called "Gold Standard" algorithm.
- Other algorithms can then be compared to it.



$$egin{aligned} oldsymbol{x'_i} &= H oldsymbol{x_i} \ oldsymbol{x'_i} &= (x_i', y_i', w_i')^T \ H oldsymbol{x_i} &= egin{bmatrix} h^{1T} oldsymbol{x_i} \ h^{2T} oldsymbol{x_i} \ h^{3T} oldsymbol{x_i} \end{bmatrix} \end{aligned}$$



$$\mathbf{x'_{i}} = H\mathbf{x_{i}}$$

$$\mathbf{x'_{i}} = (x'_{i}, y'_{i}, w'_{i})^{T}$$

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$$\mathbf{x'_{i}} = \begin{bmatrix} y'_{i}h^{3^{T}}\mathbf{x_{i}} - w'_{i}h^{2^{T}}\mathbf{x_{i}} \\ w'_{i}h^{1^{T}}\mathbf{x_{i}} - x'h^{3^{T}}\mathbf{x_{i}} \\ x'_{i}h^{2^{T}}\mathbf{x_{i}} - y'h^{1^{T}}\mathbf{x_{i}} \end{bmatrix}$$

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$$\begin{bmatrix} 0^{T} & -w'_{i}\mathbf{x_{i}}^{T} & y'_{i}\mathbf{x_{i}}^{T} \\ w'_{i}\mathbf{x_{i}}^{T} & 0^{T} & -x'_{i}\mathbf{x_{i}}^{T} \end{bmatrix}$$

$$\begin{bmatrix} h^{1} \\ h^{2} \\ h^{3} \end{bmatrix} = 0$$

- Equations are linear in  $h \to A_i h = 0$
- Only 2 out of 3 are linearly independent (indeed, 2 equations per point)

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- Holds for any homogeneous representation, e.g.  $(x_i', y_i', 1)$
- Thus holds for image points

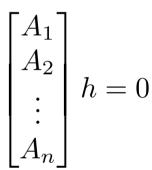
• Solving for *H* 

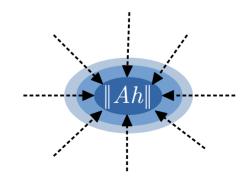
$$Ah = 0 \qquad \begin{bmatrix} A_1 \\ A_2 \\ A_3 \\ A_4 \end{bmatrix} h = 0$$

size A is 8x9 or 12x9, but rank 8

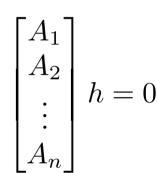
- Trivial solution is  $h=0_9^T$  is not interesting
- 1-D null-space yields solution of interest pick for example the one with: ||h||=1

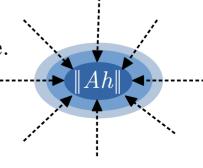
- Over-determined solution
- No exact solution because of inexact measurement, i.e. "noise".
- Find approximate solution:
  - Additional constraint needed to avoid 0, e.g. ||h||=1
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  - Additional constraint needed to avoid 0, e.g. ||h||=1
  - Since Ah=0 is not possible, so minimize ||Ah||
- The solution is the eigenvector of  $A^TA$  with the least eigenvalue.
  - That is equivalent to the singular vector corresponding to the smallest singular value of A.





#### **Objective**

Given  $n \ge 4$  2D to 2D point correspondences  $\{x_i \leftrightarrow x_i'\}$ , determine the 2D homography matrix H such that  $x_i' = Hx_i$ 

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- (iv) Determine H from h

## Some Details in the DLT



### Inhomogeneous solution

• Since h can only be computed up to scale, pick  $h_j=1$ , e.g.  $h_9=1$ , and solve for 8-vector

$$\begin{bmatrix} 0 & 0 & 0 & -x_i w_i' & -y_i w_i' & -w_i w_i' & x_i y_i' & y_i y_i' \\ x_i w_i' & y_i w_i' & w_i w_i' & 0 & 0 & 0 & x_i x_i' & y_i x_i' \end{bmatrix} \tilde{h} = \begin{bmatrix} -w_i y_i' \\ w_i x_i' \end{bmatrix}$$

- Solve using Gaussian elimination (4 points) or using linear least-squares (more than 4 points).

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- However, if  $h_9$ =0 this approach fails. Also gets poor results if  $h_9$  close to zero.

#### Therefore, not recommended.

- Note  $h_9 = H_{33} = 0$  if origin is mapped to infinity.

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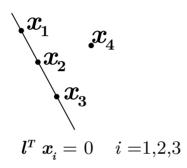
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#### Example:

When the horizon passes by the image center (0,0), that means the image center lies on the infinity!

$$m{l}_{\infty}^T H m{x}_0 = egin{bmatrix} 0 & 0 & 1 \end{bmatrix} H egin{bmatrix} 0 \ 0 \ 1 \end{bmatrix} = 0$$

Three collinear points out of 4. **PROBLEM!!!** 



Three collinear points out of 4. PROBLEM!!! 
$$x_1$$
  $x_2$   $x_3$   $x_4$   $x_2$   $x_3$   $x_4$   $x_4$   $x_2$   $x_3$   $x_4$   $x_3$   $x_4$   $x_4$   $x_5$   $x_6$   $x_$ 

Constraints: 
$$\mathbf{x}_{i}$$
,  $\mathbf{x}_{l}$ ,  $\mathbf{x}_{i} = 0$   $i = 1,2,3,4$   
Define:  $H^{*} = \mathbf{x}_{4}$ ,  $\mathbf{l}^{T}$   
Then,  $H^{*} \mathbf{x}_{i} = \mathbf{x}_{4}$ ,  $(\mathbf{l}^{T} \mathbf{x}_{i}) = 0$ ,  $i = 1,2,3$   
 $H^{*} \mathbf{x}_{4} = \mathbf{x}_{4}$ ,  $(\mathbf{l}^{T} \mathbf{x}_{4}) = k\mathbf{x}_{4}$ ,

Three collinear points out of 4.

### PROBLEM!!!

Constraints: 
$$x_i' \times Hx_i = 0$$
  $i = 1,2,3,4$ 

Define:  $H^* = x_4' l^T$ 

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Case A: If  $H^*$  is unique solution, then no homography mapping  $x_i \rightarrow x_i$  exist  $\rightarrow$  every homography must preserve collinearity.

 $H^*$  is rank-1 matrix and thus not a homography (rank-3)!

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Case B: If there is a solution H, then also  $\alpha H^* + \beta H \rightarrow$  many solutions

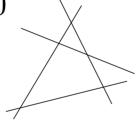
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### Solutions from lines and points

2D homographies from 2D lines

$$l'_i = H^T l_i \quad Ah = 0$$

Minimum of 4 lines

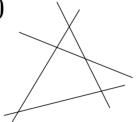


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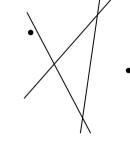
$$\mathbf{l'}_i = H^T \mathbf{l}_i \quad Ah = 0$$

Minimum of 4 lines



But can also be determined by:

- 3 general points and 1 line or
- 3 general lines and 1 point.



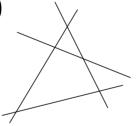
Equivalent to 4 general points

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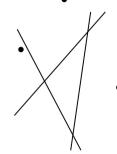
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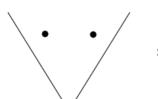
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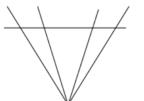


Equivalent to 4 general points

Mixed configurations that do not work. For example:

- two points and two lines = four concurrent lines = four collinear points.







# Improving DLT Normalized DLT



- Algebraic distance
- Geometric distance
- Reprojection error



# Algebraic distance

DLT minimizes 
$$||Ah|| \longrightarrow e = Ah \longrightarrow$$
 residual vector  $e_i \longrightarrow$  partial error vector for each  $(x_i \leftrightarrow x_i')$  algebraic error vector

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$$d_{alg}(\boldsymbol{x}'_i, H\boldsymbol{x}_i)^2 = \|e_i\|^2 = \left\| \begin{bmatrix} 0^T & -w_i'\boldsymbol{x}_i^T & -y_i'\boldsymbol{x}_i^T \\ -w_i'\boldsymbol{x}_i^T & 0^T & -x_i'\boldsymbol{x}_i^T \end{bmatrix} h \right\|^2 \quad algebraic \ distance$$

$$d_{alg}(\boldsymbol{x}_1, \boldsymbol{x}_2)^2 = a_1^2 + a_2^2 \text{ where } \boldsymbol{a} = (a_1, a_2, a_3)^T = \boldsymbol{x}_1 \times \boldsymbol{x}_2$$

# Algebraic distance

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$$\sum_{i} d_{alg}(\mathbf{x}'_{i}, H\mathbf{x}_{i})^{2} = \sum_{i} \|e_{i}\|^{2} = \|Ah\|^{2} = \|e\|^{2}$$

Not geometrically/statistically meaningful, but given good normalization it works fine, has an unique solution and is very fast (use for initialization)

## Geometric distance

x measured coordinates

 $\hat{x}$  estimated coordinates

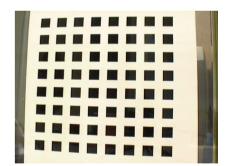
 $\bar{x}$  true coordinates

d(.,.) Euclidean distance (in image)

2 options: Consider 1 image

Consider 2 images

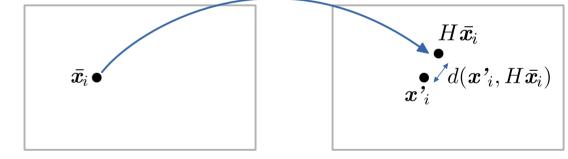
Ex: Homography between calibration pattern and its image.

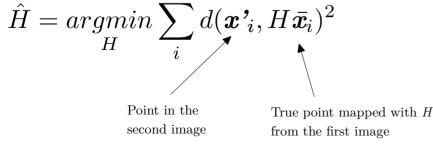


## Geometric distance

- x measured coordinates
- $\hat{x}$  estimated coordinates
- $\bar{x}$  true coordinates
- d(.,.) Euclidean distance (in image)

Error in just one image





## Geometric distance

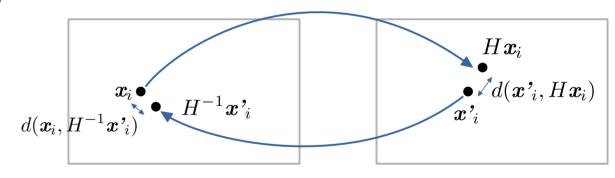
x measured coordinates

 $\hat{x}$  estimated coordinates

 $\bar{x}$  true coordinates

d(.,.) Euclidean distance (in image)

Symmetric transfer error Error in both images



$$\hat{H} = \underset{H}{\operatorname{argmin}} \sum_{i} d(\boldsymbol{x}_{i}, H^{-1}\boldsymbol{x}_{i})^{2} + d(\boldsymbol{x}_{i}, H\boldsymbol{x}_{i})^{2}$$
Transfer error in the first in the second image

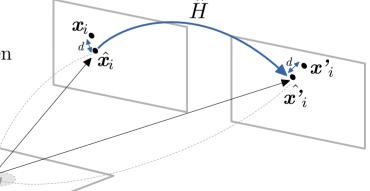
Transfer error in the second image

# Reprojection error

Minimizing the cost function involves determining the homography and a set of correspondences  $\to \hat{H}, \hat{x}_i, \hat{x}_i'$ 

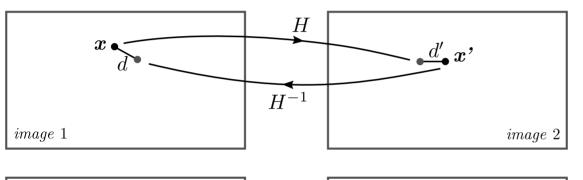
$$(\hat{H}, \hat{\boldsymbol{x}_i}, \hat{\boldsymbol{x}'_i}) = \underset{\hat{H}, \hat{\boldsymbol{x}_i}, \hat{\boldsymbol{x}'_i}}{argmin} \sum_{i} d(\boldsymbol{x}_i, \hat{\boldsymbol{x}_i})^2 + d(\boldsymbol{x}'_i, \hat{\boldsymbol{x}'_i})^2 \text{ subject to } \hat{\boldsymbol{x}'_i} = \hat{H} \hat{\boldsymbol{x}_i}$$

e.g. Estimate a point on the world plane  $\hat{X}_i$  from  $x_i \Leftrightarrow x_i$  which is then reprojected to the estimated perfectly matched correspondence  $\hat{x}_i \Leftrightarrow \hat{x}_i$ 

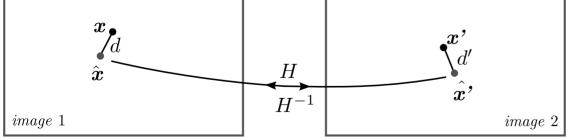


 $\hat{m{X}}_i$ 

## Symmetric transfer error X Reprojection error



$$d(x, H^{-1}x')^2 + d(x', Hx)^2$$



$$d(x, \hat{x})^2 + d(x', \hat{x'})^2$$

# Statistical cost function and Maximum Likelihood Estimation

- Optimal estimate of  $H \to \text{need a noise model}$ .
- Assume zero-mean isotropic Gaussian noise (assume outliers removed).
- Defining the pdf of the noise and the MLE of the homography results on minimizing the geometric error.
- Error in one image.

Maximum Likelihood Estimate

$$\sum_{i} d(\boldsymbol{x'_i}, H\bar{\boldsymbol{x}_i})^2$$

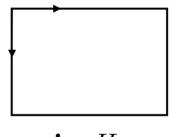
# Statistical cost function and Maximum Likelihood Estimation

- Optimal estimate of  $H \to \text{need a noise model}$ .
- Assume zero-mean isotropic Gaussian noise (assume outliers removed).
- Defining the pdf of the noise and the MLE of the homography results on minimizing the reprojection error.
- Error in both images.

Maximum Likelihood Estimate

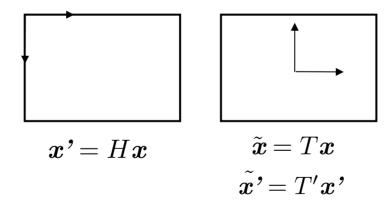
$$\sum_{i} d(\boldsymbol{x}_{i}, \hat{\boldsymbol{x}}_{i})^{2} + d(\boldsymbol{x}_{i}, \hat{\boldsymbol{x}}_{i})^{2}$$

- Are the properties and performance of the DLT algorithm invariant to transformations?
- Will result change?
- For which algorithms? For which transformations?

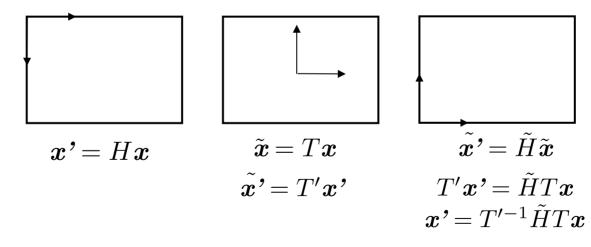


x' = Hx

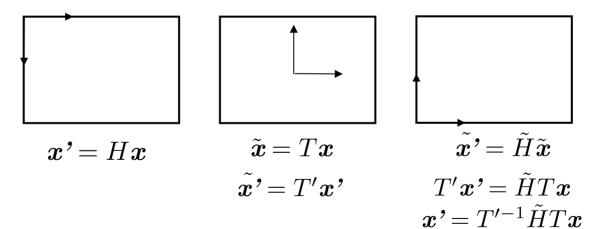
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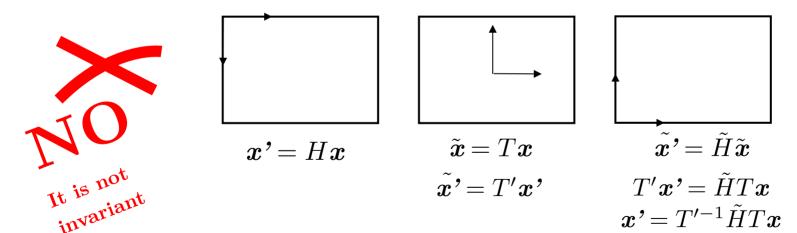


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$$? \\ H = T'^{-1}\tilde{H}T$$

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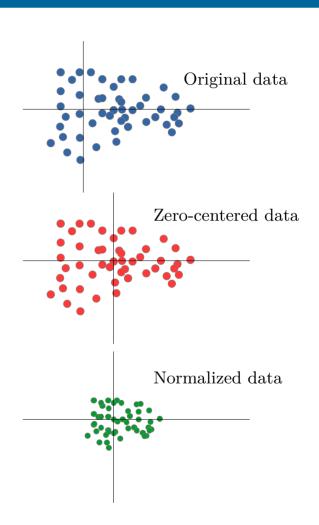
$$? \\ H = T'^{-1}\tilde{H}T$$

## Non-invariance of DLT

- DLT is not invariant to similarity transformations on the image.
- Results depend on the coordinate frame in which points are expressed.
- Some coordinate systems are in some way better than others for computing a 2D homography.
- Data normalization is an essential step in the DLT algorithm. It **MUST NOT** be considered optional.

# Normalizing transformations

- Algorithms with initial normalization step will be invariant to arbitrary choices of scale and coordinate origin
- What is a good choice of coordinates?
  - Translate centroid to origin.
  - Scale to a  $\sqrt{2}$  average distance to the origin. This means that the average point is equal to  $(1,1,1)^T$ .
  - This transformation is applied independently on both images.



# Importance of normalization

$$\begin{bmatrix} 0 & 0 & 0 & -\boldsymbol{x'_i} & -\boldsymbol{y'_i} & -1 & \boldsymbol{y'_i}\boldsymbol{x_i} & \boldsymbol{y'_i}\boldsymbol{y_i} & \boldsymbol{y'_i} \\ \boldsymbol{x_i} & \boldsymbol{y_i} & 1 & 0 & 0 & 0 & -\boldsymbol{x'_i}\boldsymbol{x_i} & -\boldsymbol{x'_i}\boldsymbol{y_i} & -\boldsymbol{x'_i} \end{bmatrix} \begin{bmatrix} h^1 \\ h^2 \\ h^3 \end{bmatrix} = 0$$

$$\sim 10^2 \sim 10^2 \ 1 \sim 10^2 \ \sim 10^2 \ 1 \sim 10^4 \ \sim 10^4 \ \sim 10^4$$

#### orders of magnitude difference!

The effect of normalization (simulation): 5 point (crosses) were used to compute a 2D Homography. The homography H is the identity mapping. 100 trials were made adding 0.1 pixel Gaussian noise to the points. The result H was applied to a further point. (a) result without normalization and (b) with normalization.

#### **Objective**

Given  $n \ge 4$  2D to 2D point correspondences  $\{x_i \leftrightarrow x_i'\}$ , determine the 2D homography matrix H such that  $x_i' = Hx_i$ 

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#### **Objective**

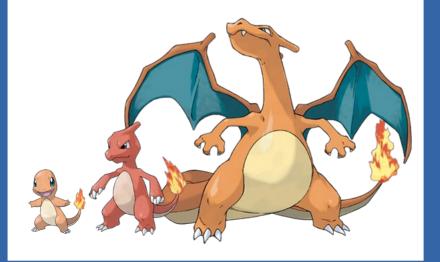
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#### Algorithm

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- Apply DLT algorithm to  $\tilde{x}_i \leftrightarrow \tilde{x}_i$ '
- Denormalize solution  $H = T'^{-1}\tilde{H}T$

# Normalized DLT





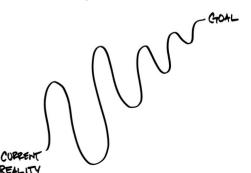
# Iterative minimization methods

Required to minimize geometric error:		Therefore, careful implementation required:	
(i)	Often slower than DLT	(i)	Cost function
(ii)	Require initialization	(ii)	Parameterization (minimal or not)
(iii)	No guaranteed convergence, local minima	(iii)	Function specification(cost $\leftrightarrow$ parameters)
(iv)	Stopping criterion required	(iv)	Initialization
		(v)	Iterations

## Parametrization

- Parameters should cover complete space and allow efficient estimation of cost.
- Minimal or over-parameterized? e.g. 8 or 9
  - Minimal often more complex, also cost surface.
  - Sometimes stuck in local minimum.
  - Good algorithms can deal with over-parameterization.
- Parametrization can also be used to restrict transformation to particular class.





# Function specifications

- Measurement vector  $X \in \mathbb{R}^N$  with covariance  $\Sigma$
- (i) Set of parameters represented by vector  $P \in \mathbb{R}^{M}$
- (ii) Mapping  $f: \mathbb{R}^{M} \to \mathbb{R}^{N}$
- (iii) Cost function to be minimized

The goal is to find a set of parameters P such that f(P) = X or failing that, to bring f(P) as close to X as possible

#### Error in one image

- Points in the first image without error.
- Measurements  $\rightarrow$  point in the second image.
- Find *H* that minimizes the cost function.

$$\sum_{i} d(\boldsymbol{x}_{i}^{\prime}, H\bar{\boldsymbol{x}}_{i})^{2}$$

$$f: h \to (H\bar{\boldsymbol{x}}_{1}, H\bar{\boldsymbol{x}}_{2}, \dots, H\bar{\boldsymbol{x}}_{n})$$

$$\|\boldsymbol{X} - f(h)\|$$

#### Error in one image

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## Symmetric transfer error

- Measurements  $\rightarrow$  points in both images.
- Find H that minimizes the cost function.

$$\sum_{i} d(\boldsymbol{x}_{i}^{\prime}, H\bar{\boldsymbol{x}}_{i})^{2}$$

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$$\sum_{i} d(\boldsymbol{x}_{i}, H^{-1}\boldsymbol{x}_{i}^{\prime})^{2} + d(\boldsymbol{x}_{i}^{\prime}, H\boldsymbol{x}_{i})^{2}$$

$$f: h \to (H^{-1}\boldsymbol{x}_{1}^{\prime}, \dots, H^{-1}\boldsymbol{x}_{n}^{\prime}, H\boldsymbol{x}_{1}, \dots, H\boldsymbol{x}_{n})$$

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## Symmetric transfer error

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#### Reprojection error

• Find H and points that minimizes the cost function.

$$\sum_{i} d(\boldsymbol{x'_i}, H\bar{\boldsymbol{x}_i})^2$$
 $f: h \to (H\boldsymbol{x}_1, H\boldsymbol{x}_2, \dots, H\boldsymbol{x}_n)$ 
 $\|\boldsymbol{X} - f(h)\|$ 

$$\sum_{i} d(\boldsymbol{x}_{i}, H^{-1}\boldsymbol{x}_{i}^{\prime})^{2} + d(\boldsymbol{x}_{i}^{\prime}, H\boldsymbol{x}_{i})^{2}$$

$$f: h \to (H^{-1}\boldsymbol{x}_{1}^{\prime}, \dots, H^{-1}\boldsymbol{x}_{n}^{\prime}, H\boldsymbol{x}_{1}, \dots, H\boldsymbol{x}_{n})$$

$$\|\boldsymbol{X} - f(h)\|$$

$$\sum_{i} d(\boldsymbol{x}_{i}, \hat{\boldsymbol{x}}_{i})^{2} + d(\boldsymbol{x}_{i}, \hat{\boldsymbol{x}}_{i})^{2}$$

$$f: (h, \hat{\boldsymbol{x}}_{1}, \dots, \hat{\boldsymbol{x}}_{n}) \rightarrow (\hat{\boldsymbol{x}}_{1}, \hat{\boldsymbol{x}}_{1}, \dots, \hat{\boldsymbol{x}}_{n}, \hat{\boldsymbol{x}}_{n})$$

$$\|\boldsymbol{X} - f(h)\|$$

## Initialization

- Typically, use linear solution
- If outliers, use robust algorithm

## Iteration methods

- Most popular:
  - Newton's method
  - Levenberg-Marquardt





# Gold Standard algorithm

#### **Objective**

Given  $n \ge 4$  2D to 2D point correspondences  $\{x_i \longleftrightarrow x_i'\}$ , determine the Maximum Likelyhood Estimation of H (this also implies computing optimal  $\hat{x}_i' = \hat{H}\hat{x}_i$ )

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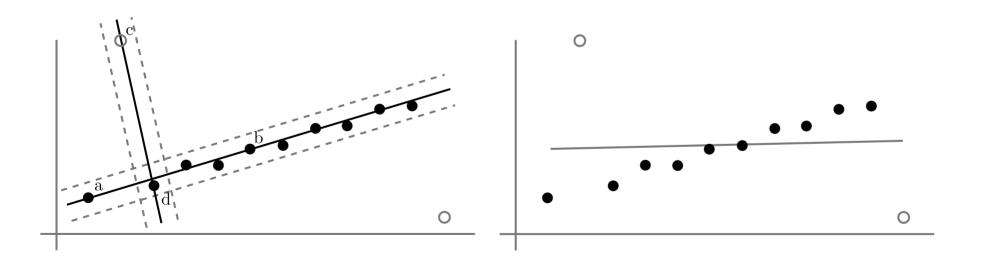
- (i) Initialization: compute an initial estimate using normalized DLT or RANSAC
- (ii) Geometric minimization of Gold Standard error:
  - compute initial estimate for optimal  $\{\hat{x}_i\}$  from  $\{x_i\}$
  - minimize cost  $\Sigma d(\boldsymbol{x}_i, \hat{\boldsymbol{x}}_i)^2 + d(\boldsymbol{x}_i', \hat{\boldsymbol{x}}_i')^2$  over  $\{\hat{H}, \hat{\boldsymbol{x}}_1, \hat{\boldsymbol{x}}_2, \hat{\boldsymbol{x}}_3, ..., \hat{\boldsymbol{x}}_n\}$
  - if many points, use sparse method

# RANSAC RANdom Sample Consensus



## Robust estimation

• What if set of matches contains gross outliers?



## **Objective**

Robust fit of model to data set S which contains outliers

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(i) Randomly select a sample of s data points from S and instantiate the model from this subset.

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- (iv) If the size of  $S_i$  is less than T, select a new subset and repeat the above.
- (v) After N trials the largest consensus set  $S_i$  is selected, and the model is re-estimated using all the points in the subset  $S_i$

## Distance threshold

Choose t so probability for inlier is  $\alpha$  (e.g. 0.95)

- Often empirically.
- Zero-mean Gaussian noise  $\sigma$  then the square of the point distance  $d_{\perp}^2$  follows a  $X_m^2$  distribution with m = codimension of model.

 $(dimension+codimension=dimension\ space)$ 

Codimension	Model	$t^{2}$
1	$_{ m l,F}$	$3.84  \sigma^2$
2	$_{\mathrm{H,P}}$	$5.99  \sigma^2$
3	${ m T}$	$7.81  \sigma^2$

# How many samples?

Choose N so that, with probability p, at least one random sample is free from outliers. e.g. p=0.99. Suppose e is the propability that any selected point is an outlier, then at least N selections (each of s points) are require so that  $(1-(1-e)^s)^N=1-p$ .

Sample size	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

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

# Acceptable consensus set?

• Typically, terminate when the consensus set is similar to the expected number of inliers

$$T = (1 - e)n$$

- Adaptively determining the number of samples:
  - e is often unknown a priori, so pick worst case, e.g. 50%, and adapt if more inliers are found, e.g. 80% would yield e=0.2
  - $N=\infty$ , sample count =0
  - While  $N > sample\_count$  repeat
    - Choose a sample and count the number of inliers
    - Set e=1-(number of inliers)/(total number of points)
    - Recompute N from e, usually with p = 0.99
    - Increment the sample\_count by 1

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

# Robust Algorithm

• Use RANSAC to maximize the number of inliers



# Robust Algorithm

• Use RANSAC to maximize the number of inliers

Bad matches  $\rightarrow$  outliers



#### <u>Objective</u>

Compute homography between two images

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#### Algorithm

(i) Interest points: Compute interest points in each image

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- (i) Interest points: Compute interest points in each image
- (ii) Putative correspondences: Compute a set of interest point matches based on some similarity measure
- (iii) RANSAC robust estimation: Repeat for N samples
  - (a) Select 4 correspondences and compute H
  - (b) Calculate the distance  $d_i$  for each putative match
  - (c) Compute the number of inliers consistent with H ( $d_i < t$ )

Choose H with most inliers

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- (vii) Optimal estimation: re-estimate H from all inliers by minimizing ML cost function with Levenberg-Marquardt
- (viii) Guided matching: Determine more matches using prediction by computed H

Optionally iterate last two steps until convergence

# Example

Image 1

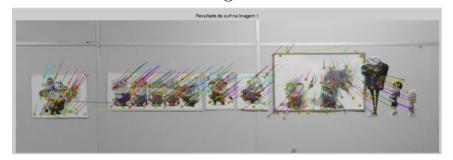


Image  $2 \to \text{Image } 1$ 



Image 2



Image 1  $\rightarrow$  Image 2



## Determine putative correspondences

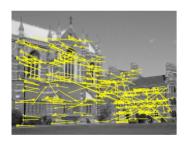
- Compare interest points using similarity measure:
  - SAD, SSD, ZNCC on small neighborhood
- If motion is limited, only consider interest points with similar coordinates
- More advanced approaches exist, based on invariance...
  - SIFT
  - SURF
  - others

## Example: robust computation





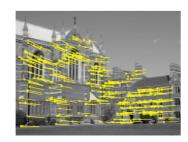
Interest points (500/image)





Putative correspondences (268) Outliers (117)





Inliers (151) Final inliers (262)

## Credits

- Richard Hartley and Andrew Zisserman. Multiple View
   Geometry in Computer Vision. Cambridge,
   ISBN 0521623049
- Based on slides from Marc Pollefeys https://www.cs.unc.edu/~marc/mvg/slides.html