

# Computer Vision: Image Alignment

Raquel Urtasun

TTI Chicago

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- Chapter 2.1, 3.6, 4.3 and 6.1 of Szeliski's book
- Chapter 1 of Forsyth & Ponce

What did we see in class last week?

# What is the geometric relationship between these images?



[Source: N. Snavely]

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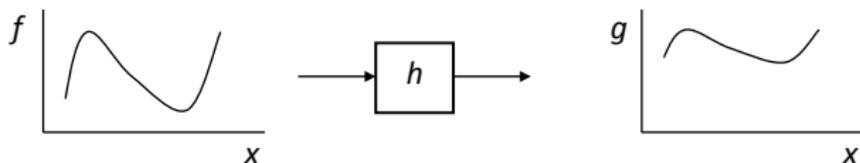
Very important for creating mosaics!

[Source: N. Snavely]

# Image Warping

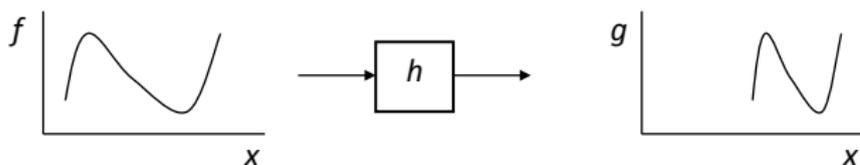
- **Image filtering:** change *range* of image

$$g(x) = h(f(x))$$



- **Image warping:** change *domain* of image

$$g(x) = f(h(x))$$



[Source: R. Szeliski]

# Parametric (global) warping



$\mathbf{p} = (x, y)$



$\mathbf{p}' = (x', y')$

- Transformation  $T$  is a coordinate-changing machine:

$$p' = T(p)$$

- What does it mean that  $T$  is global?
  - Is the same for any point  $p$
  - Can be described by just a few numbers (parameters)

[Source: N. Snavely]

# Forward and Inverse Warping

- **Forward Warping:** Send each pixel  $f(x)$  to its corresponding location  $(x', y') = T(x, y)$  in  $g(x', y')$

```
procedure forwardWarp( $f, h$ , out  $g$ ):
```

```
  For every pixel  $x$  in  $f(x)$ 
```

1. Compute the destination location  $x' = h(x)$ .
2. Copy the pixel  $f(x)$  to  $g(x')$ .

- **Inverse Warping:** Each pixel at the destination is sampled from the original image

```
procedure inverseWarp( $f, h$ , out  $g$ ):
```

```
  For every pixel  $x'$  in  $g(x')$ 
```

1. Compute the source location  $x = \hat{h}(x')$
2. Resample  $f(x)$  at location  $x$  and copy to  $g(x')$

# All 2D Linear Transformations

Linear transformations are combinations of

- Scale,
- Rotation
- Shear
- Mirror

$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} a & b \\ c & d \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix}$$

[Source: N. Snavely]

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Properties of linear transformations:

- Origin maps to origin
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$$\begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} a & b \\ c & d \end{bmatrix} \begin{bmatrix} e & f \\ g & h \end{bmatrix} \begin{bmatrix} i & j \\ k & l \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix}$$

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# Affine Transformations

Affine transformations are combinations of

- Linear transformations, and
- Translations

$$\begin{bmatrix} x' \\ y' \\ w \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} x \\ y \\ w \end{bmatrix}$$

Properties of affine transformations:

- Origin does not necessarily map to origin
- Lines map to lines
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[Source: N. Snavely]

# Projective Transformations

- Affine transformations and Projective warps

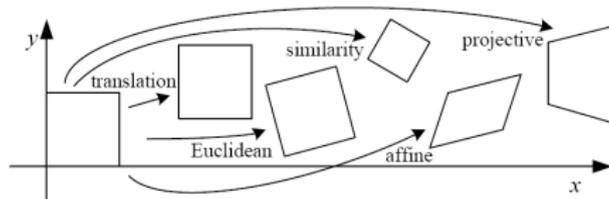
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[Source: N. Snavely]

# 2D Image Transformations



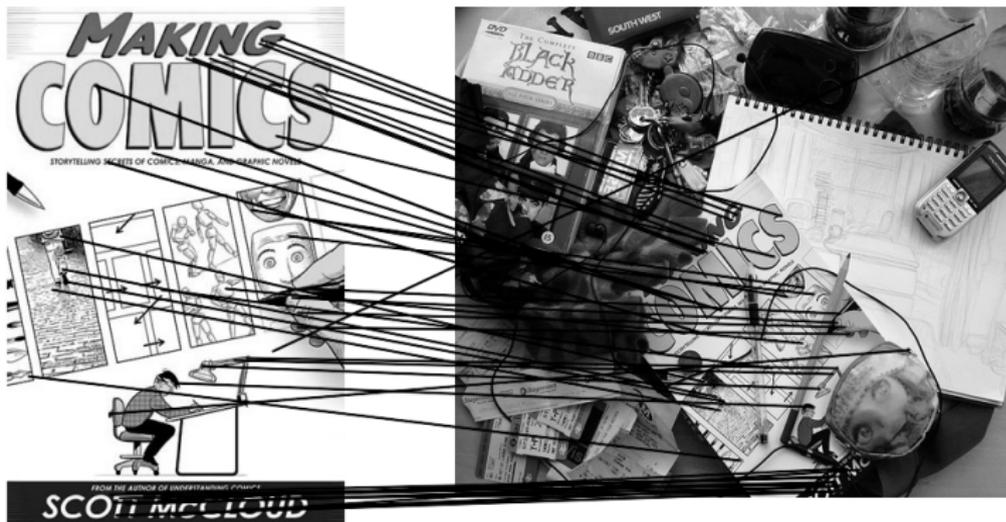
Transformation	Matrix	# DoF	Preserves	Icon
translation	$\begin{bmatrix} \mathbf{I} & \mathbf{t} \end{bmatrix}_{2 \times 3}$	2	orientation	
rigid (Euclidean)	$\begin{bmatrix} \mathbf{R} & \mathbf{t} \end{bmatrix}_{2 \times 3}$	3	lengths	
similarity	$\begin{bmatrix} s\mathbf{R} & \mathbf{t} \end{bmatrix}_{2 \times 3}$	4	angles	
affine	$\begin{bmatrix} \mathbf{A} \end{bmatrix}_{2 \times 3}$	6	parallelism	
projective	$\begin{bmatrix} \tilde{\mathbf{H}} \end{bmatrix}_{3 \times 3}$	8	straight lines	

- These transformations are a nested set of groups
- Closed under composition and inverse is a member

# Computing transformations

Given a set of matches between images A and B

- How can we compute the transform  $T$  from A to B?
- Find transform  $T$  that best agrees with the matches



[Source: N. Snavely]

# Least squares formulation

- For each point  $(x_i, y_i)$  we have

$$\begin{aligned}x_i + x_t &= x'_i \\y_i + y_t &= y'_i\end{aligned}$$

- We define the residuals as

$$\begin{aligned}r_{x_i}(x_t) &= x_i + x_t - x'_i \\r_{y_i}(y_t) &= y_i + y_t - y'_i\end{aligned}$$

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- **Goal:** minimize sum of squared residuals

$$C(x_t, y_t) = \sum_{i=1}^n (r_{x_i}(x_t)^2 + r_{y_i}(y_t)^2)$$

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[Source: N. Snavely]

- We can also write as a matrix equation

$$\begin{bmatrix} 1 & 0 \\ 0 & 1 \\ 1 & 0 \\ 0 & 1 \\ \vdots & \\ 1 & 0 \\ 0 & 1 \end{bmatrix} \begin{bmatrix} x_t \\ y_t \end{bmatrix} = \begin{bmatrix} x'_1 - x_1 \\ y'_1 - y_1 \\ x'_2 - x_2 \\ y'_2 - y_2 \\ \vdots \\ x'_n - x_n \\ y'_n - y_n \end{bmatrix}$$
$$\mathbf{A} \quad \mathbf{t} = \quad \mathbf{b}$$

$2n \times 2$        $2 \times 1$        $2n \times 1$

- Solve for  $\mathbf{t}$  by looking at the fixed-point equation

# Affine Transformations

When we are dealing with an affine transformation

$$\begin{bmatrix} x' \\ y' \\ w' \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} x \\ y \\ w \end{bmatrix}$$

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# Affine Transformation Cost Function

- We can write the residuals as

$$\begin{aligned}r_{x_i}(a, b, c, d, e, f) &= (ax_i + by_i + c) - x'_i \\r_{y_i}(a, b, c, d, e, f) &= (dx_i + ey_i + f) - y'_i\end{aligned}$$

- Cost function

$$C(a, b, c, d, e, f) = \sum_{i=1}^N (r_{x_i}(a, b, c, d, e, f)^2 + r_{y_i}(a, b, c, d, e, f)^2)$$

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# Matrix form

$$\begin{bmatrix} x_1 & y_1 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & x_1 & y_1 & 1 \\ x_2 & y_2 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & x_2 & y_2 & 1 \\ & & \vdots & & & \\ x_n & y_n & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & x_n & y_n & 1 \end{bmatrix} \begin{bmatrix} a \\ b \\ c \\ d \\ e \\ f \end{bmatrix} = \begin{bmatrix} x'_1 \\ y'_1 \\ x'_2 \\ y'_2 \\ \vdots \\ x'_n \\ y'_n \end{bmatrix}$$
$$\mathbf{A}_{2n \times 6} \mathbf{t}_{6 \times 1} = \mathbf{b}_{2n \times 1}$$

[Source: N. Snavely]

- Let  $x' = f(x; p)$  be a parametric transformation
- In the case of translation, similarity and affine, there is a linear relationship between the amount of motion  $\Delta x = x' - x$  and the unknown parameters

$$\Delta x = x' - x = \mathbf{J}(x)\mathbf{p}$$

with  $\mathbf{J} = \frac{\partial f}{\partial p}$  is the **Jacobian** of the transformation  $\mathbf{f}$  with respect to the motion parameters  $\mathbf{p}$

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# General Formulation

Transform	Matrix	Parameters $p$	Jacobian $J$
translation	$\begin{bmatrix} 1 & 0 & t_x \\ 0 & 1 & t_y \end{bmatrix}$	$(t_x, t_y)$	$\begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}$
Euclidean	$\begin{bmatrix} c_\theta & -s_\theta & t_x \\ s_\theta & c_\theta & t_y \end{bmatrix}$	$(t_x, t_y, \theta)$	$\begin{bmatrix} 1 & 0 & -s_\theta x - c_\theta y \\ 0 & 1 & c_\theta x - s_\theta y \end{bmatrix}$
similarity	$\begin{bmatrix} 1 + a & -b & t_x \\ b & 1 + a & t_y \end{bmatrix}$	$(t_x, t_y, a, b)$	$\begin{bmatrix} 1 & 0 & x & -y \\ 0 & 1 & y & x \end{bmatrix}$
affine	$\begin{bmatrix} 1 + a_{00} & a_{01} & t_x \\ a_{10} & 1 + a_{11} & t_y \end{bmatrix}$	$(t_x, t_y, a_{00}, a_{01}, a_{10}, a_{11})$	$\begin{bmatrix} 1 & 0 & x & y & 0 & 0 \\ 0 & 1 & 0 & 0 & x & y \end{bmatrix}$

- Let's do a couple on the board!

- The sum of square residuals is then

$$\begin{aligned} E_{LLS} &= \sum_i \|\mathbf{J}(\mathbf{x}_i)\mathbf{p} - \Delta\mathbf{x}_i\|_2^2 \\ &= \mathbf{p}^T \left[ \sum_i \mathbf{J}^T(\mathbf{x}_i)\mathbf{J}(\mathbf{x}_i) \right] \mathbf{p} - 2\mathbf{p}^T \left[ \sum_i \mathbf{J}^T(\mathbf{x}_i)\Delta\mathbf{x}_i \right] + \sum_i \|\Delta\mathbf{x}_i\|_2^2 \end{aligned}$$

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- We can compute the solution by looking for a fixed point, yielding

$$\mathbf{A}\mathbf{p} = \mathbf{b}$$

with  $\mathbf{A} = \sum_i \mathbf{J}^T(\mathbf{x}_i)\mathbf{J}(\mathbf{x}_i)$  the **Hessian** and  $\mathbf{b} = \sum_i \mathbf{J}^T(\mathbf{x}_i)\Delta\mathbf{x}_i$

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# Uncertainty Weighting

- The above solution assumes that all feature points are matched with same accuracy.
- If we associate a scalar variance  $\sigma_i^2$  with each correspondence, we can minimize the **weighted least squares** problem

$$E_{WLS} = \sum_i \sigma_i^{-2} \|\mathbf{r}_i\|_2^2$$

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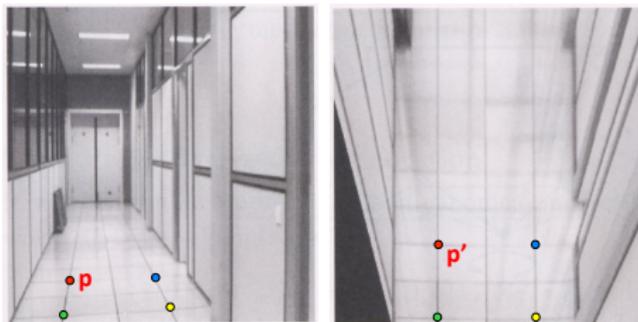
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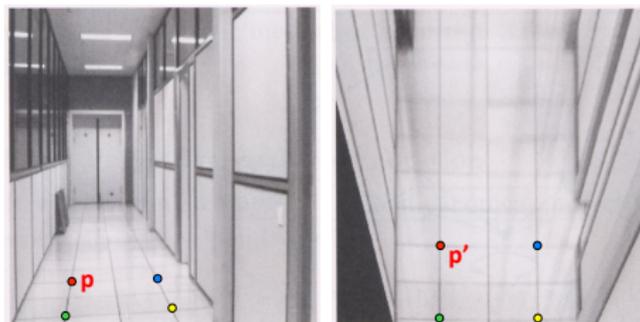
# Homographies



To unwarp (rectify) and image

- solve for homography  $H$  given  $p$  and  $p'$

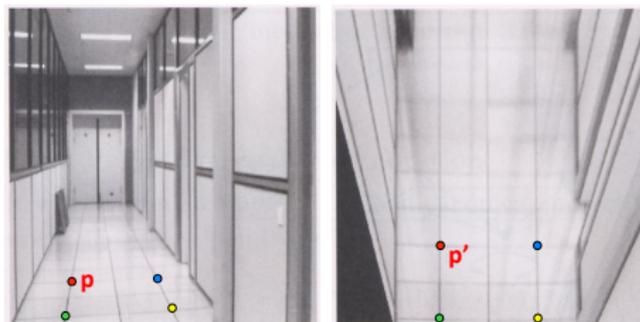
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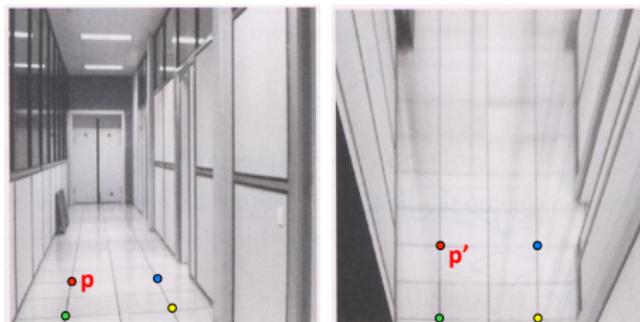
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  - linear in unknowns:  $\mathbf{w}$  and coefficients of  $\mathbf{H}$

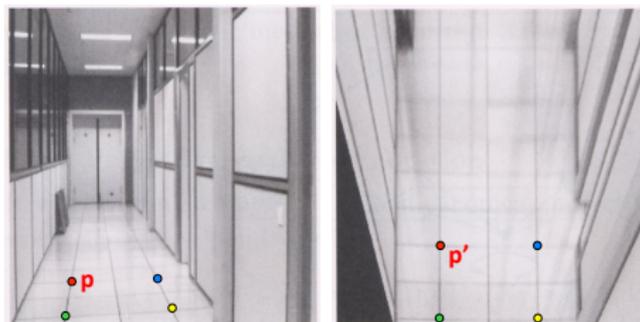
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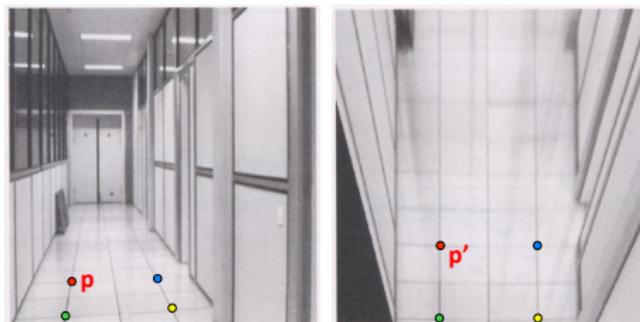


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  - linear in unknowns:  $\mathbf{w}$  and coefficients of  $\mathbf{H}$
  - $\mathbf{H}$  is defined up to an arbitrary scale factor
  - how many points are necessary to solve for  $\mathbf{H}$ ?

[Source: N. Snavely]

# Homographies



To unwarp (rectify) and image

- solve for homography  $H$  given  $p$  and  $p'$
- solve equations of the form:  $\mathbf{w}p' = \mathbf{H}p$ 
  - linear in unknowns:  $\mathbf{w}$  and coefficients of  $\mathbf{H}$
  - $\mathbf{H}$  is defined up to an arbitrary scale factor
  - how many points are necessary to solve for  $\mathbf{H}$ ?

[Source: N. Snavely]

# Solving for Homographies

$$\begin{bmatrix} ax'_i \\ ay'_i \\ a \end{bmatrix} = \begin{bmatrix} h_{00} & h_{01} & h_{02} \\ h_{10} & h_{11} & h_{12} \\ h_{20} & h_{21} & h_{22} \end{bmatrix} \begin{bmatrix} x_i \\ y_i \\ 1 \end{bmatrix}$$

- To get to non-homogenous coordinates

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- This is still linear in the unknowns

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$\mathbf{A}$   $\mathbf{h}$   $\mathbf{0}$

$2n \times 9$   $9$   $2n$

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$$\min_{\mathbf{h}} \|\mathbf{Ah}\|_2^2$$

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Given images  $A$  and  $B$

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Is there a problem with this?

[Source: N. Snavely]

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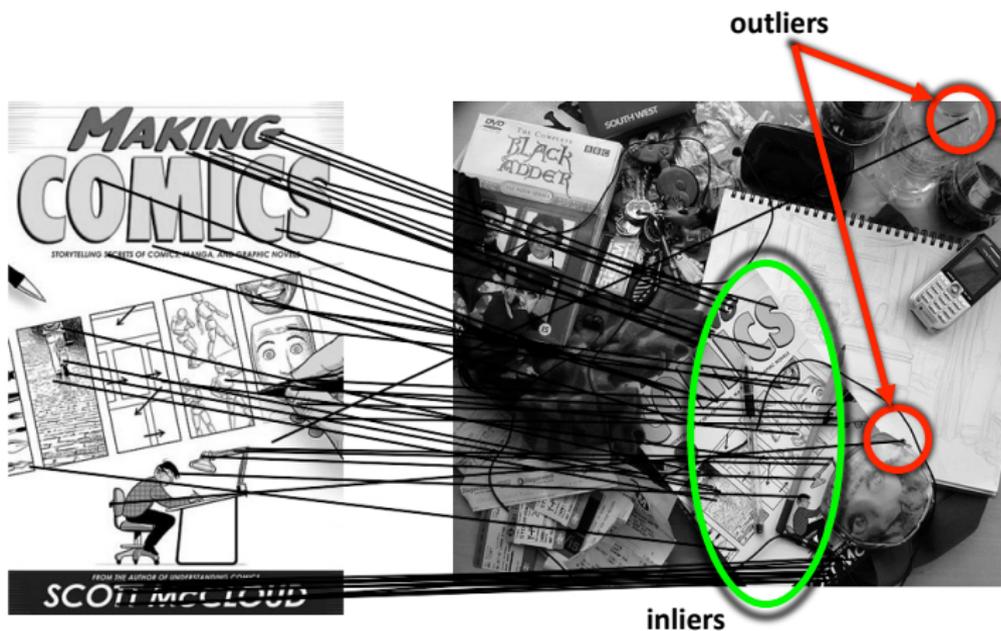
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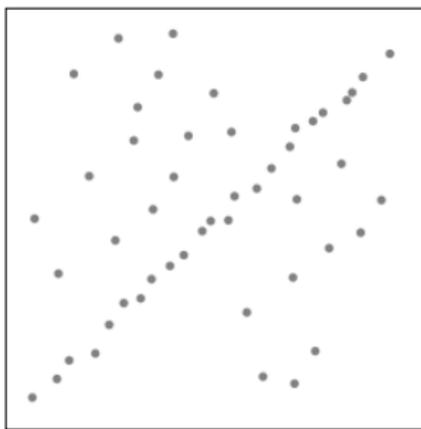
# Robustness



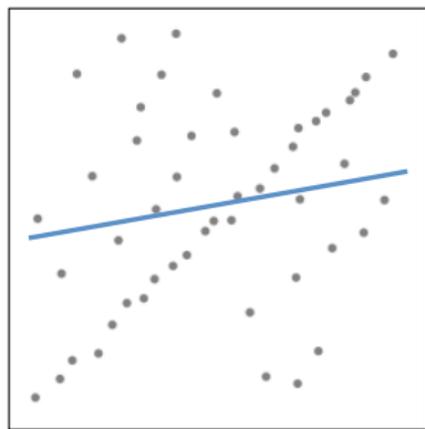
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- Lets consider a simpler example ... linear regression



Problem: Fit a line to these datapoints

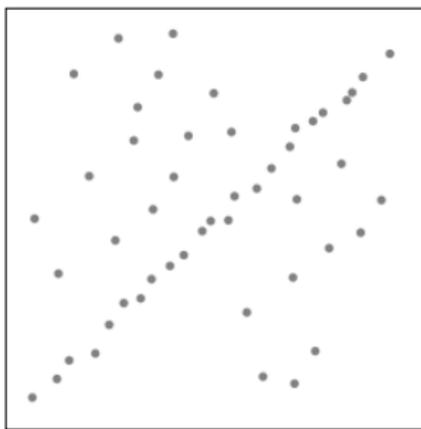


Least squares fit

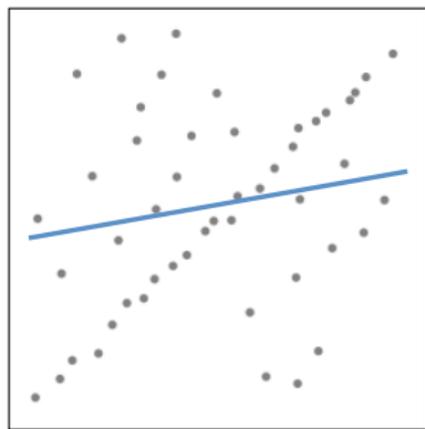
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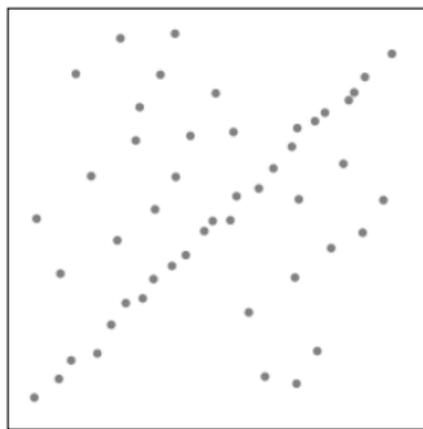
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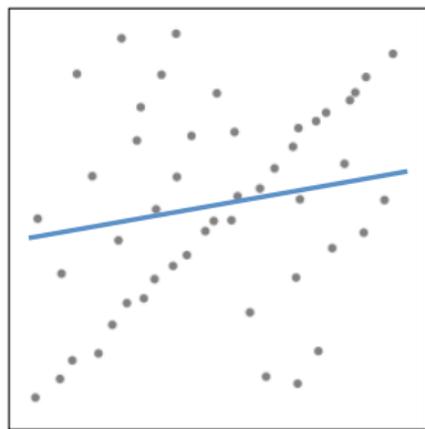
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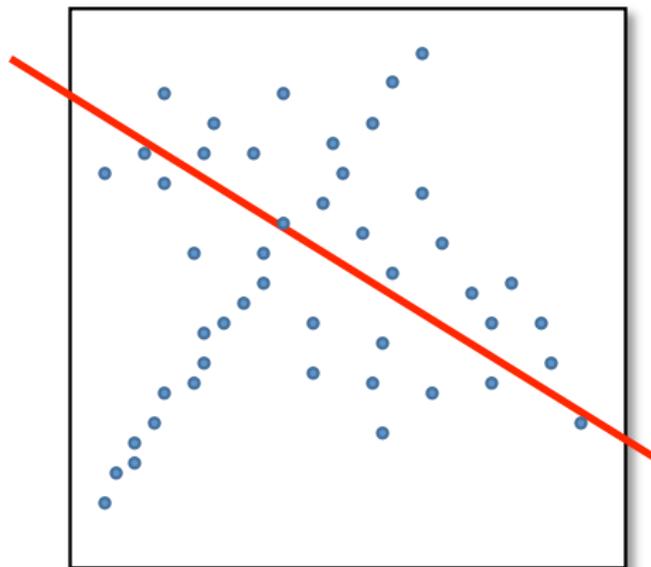
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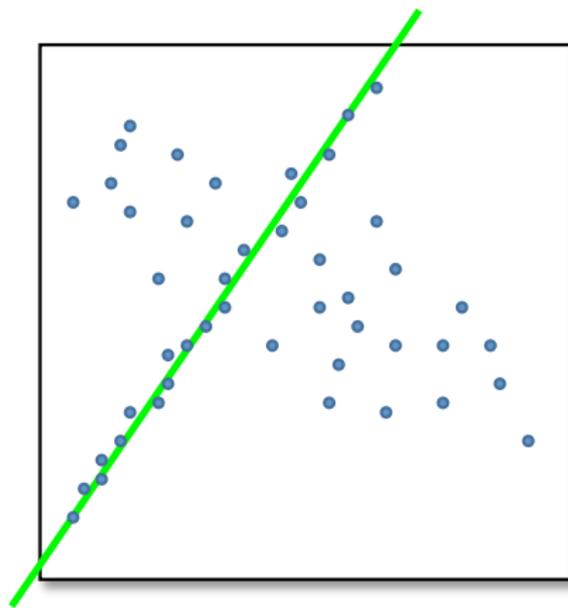
# Counting Inliers



**Inliers: 3**

[Source: N. Snavely]

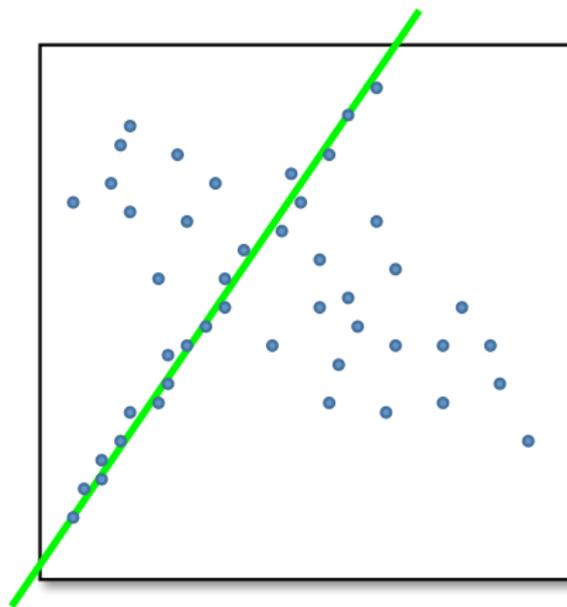
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**Inliers: 20**

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What's the problem with this approach?

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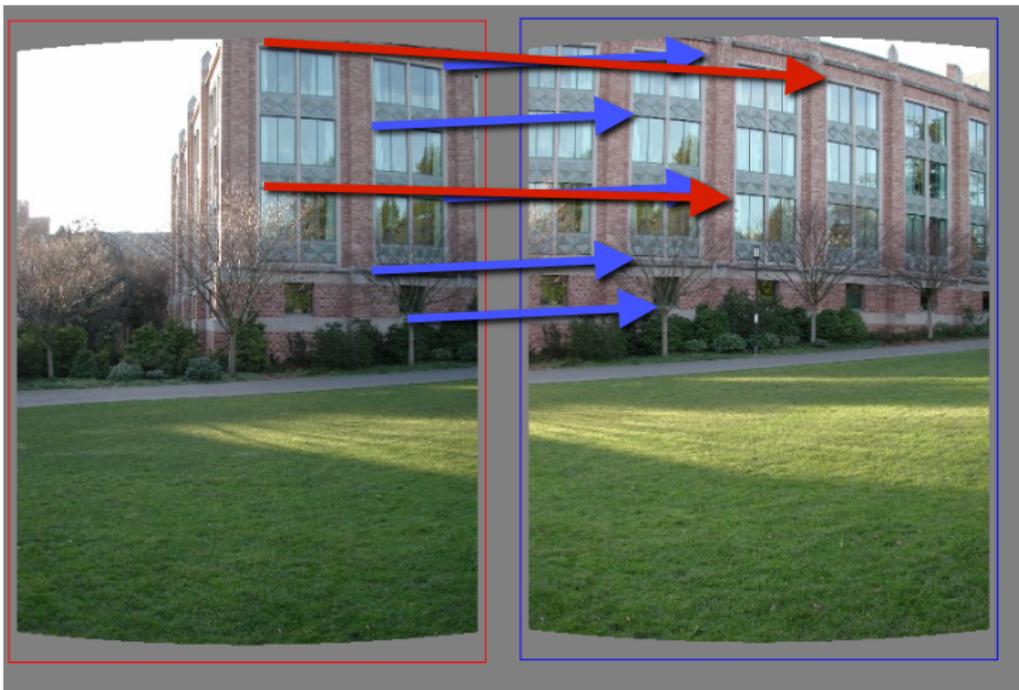
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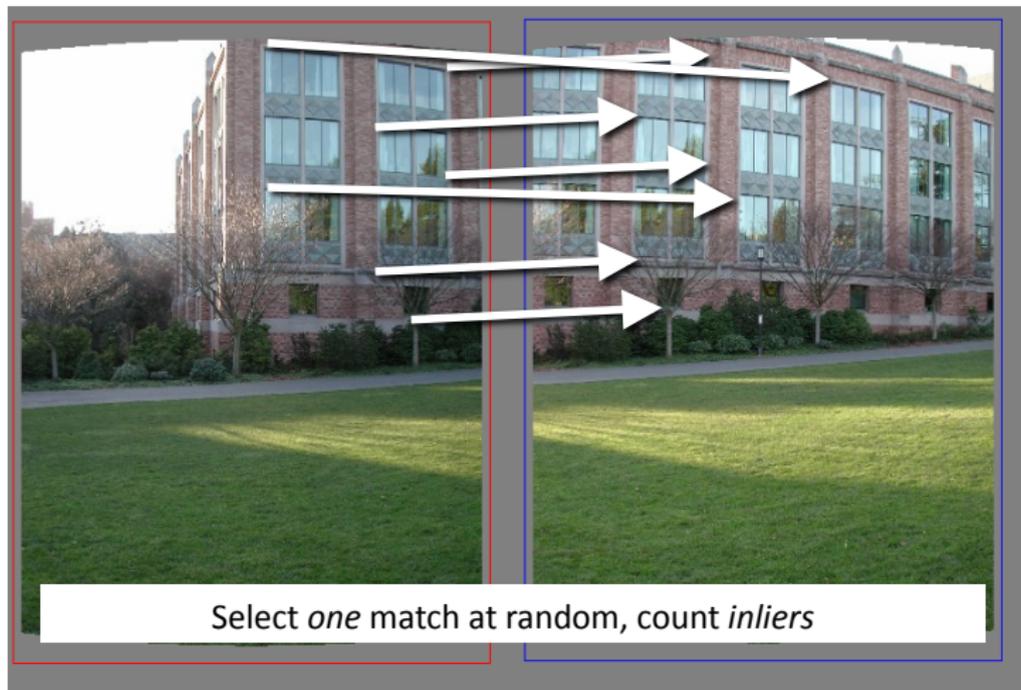
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# Translations



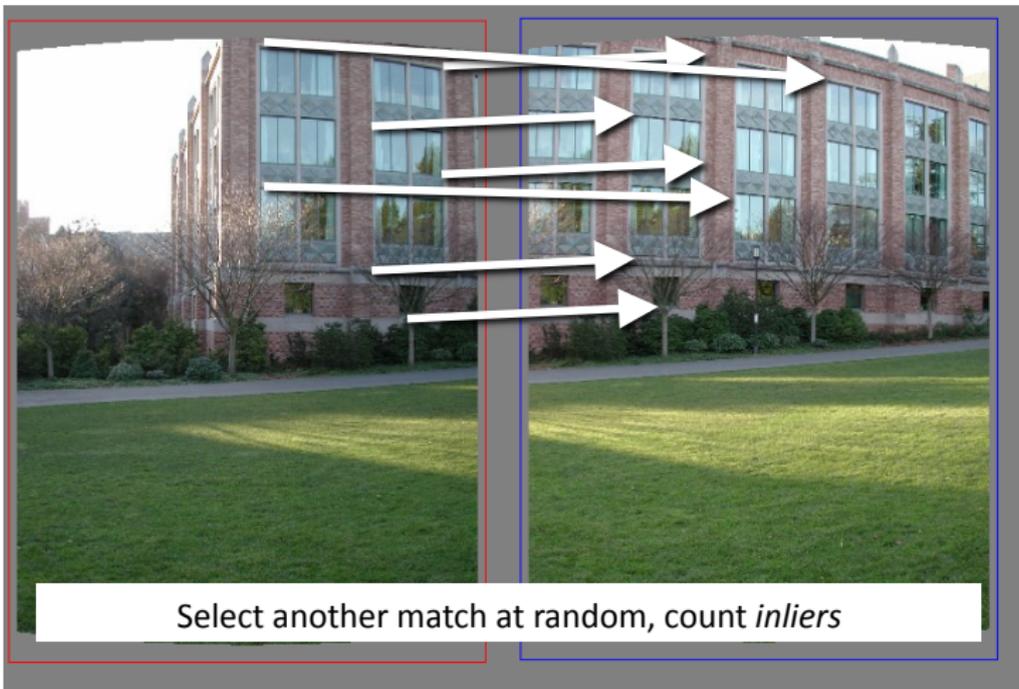
[Source: N. Snavely]

# RANdom SAmple Consensus



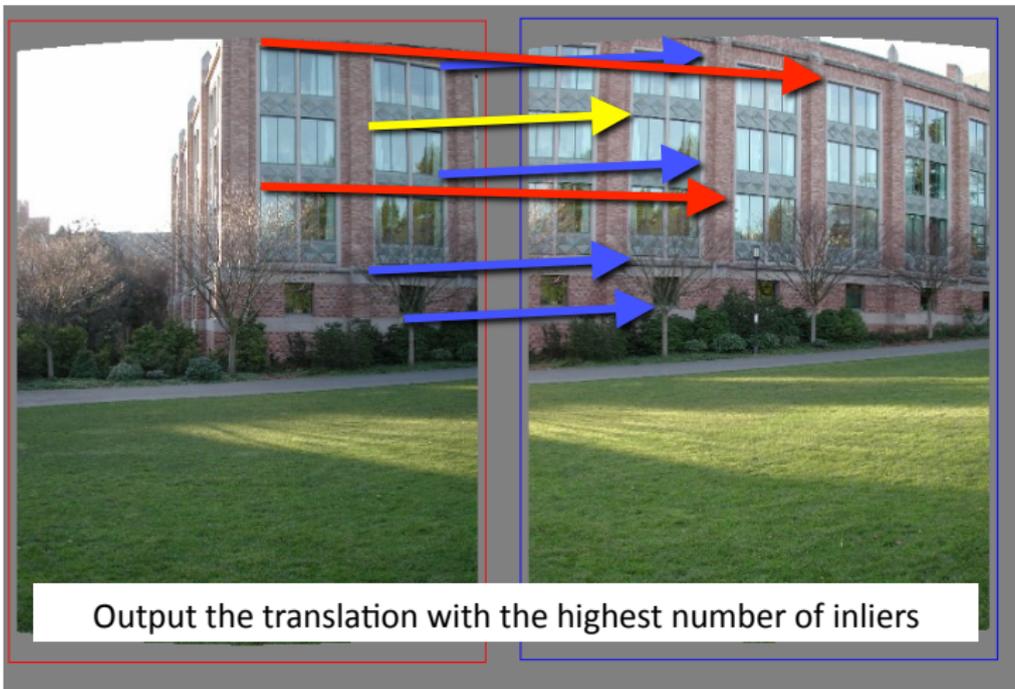
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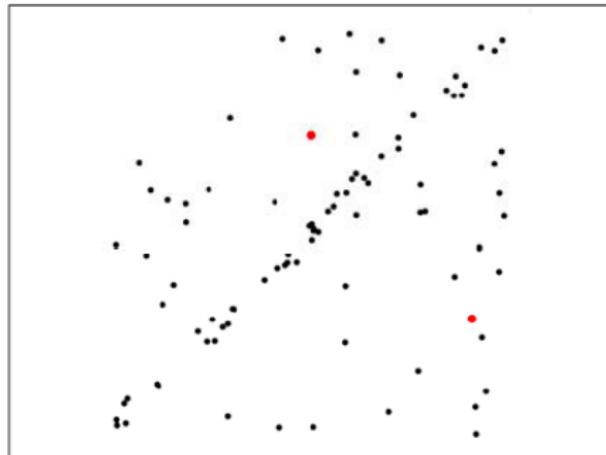
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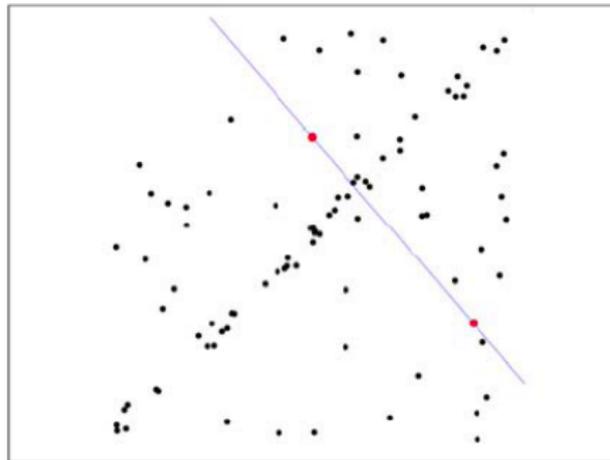
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[Source: R. Raguram]

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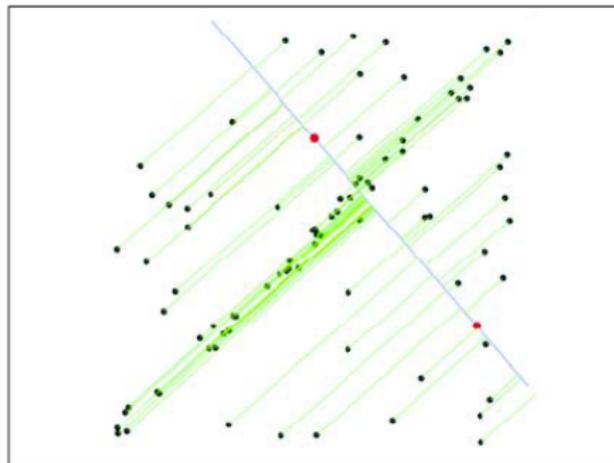
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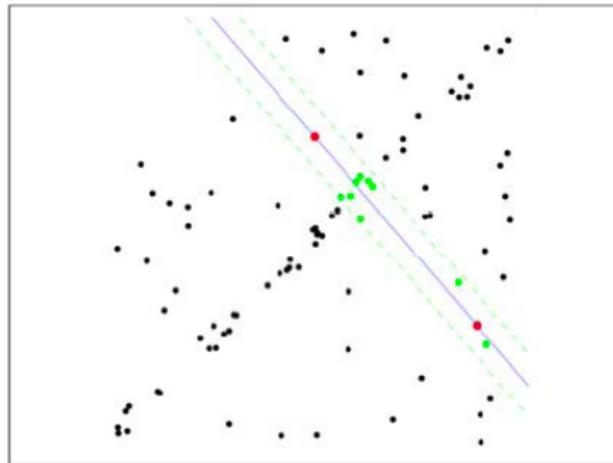
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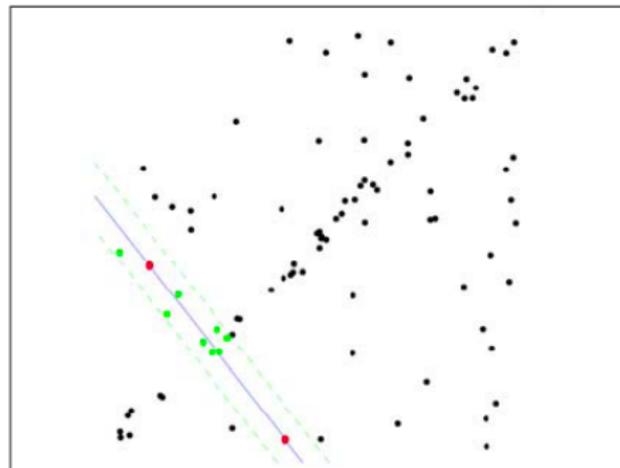
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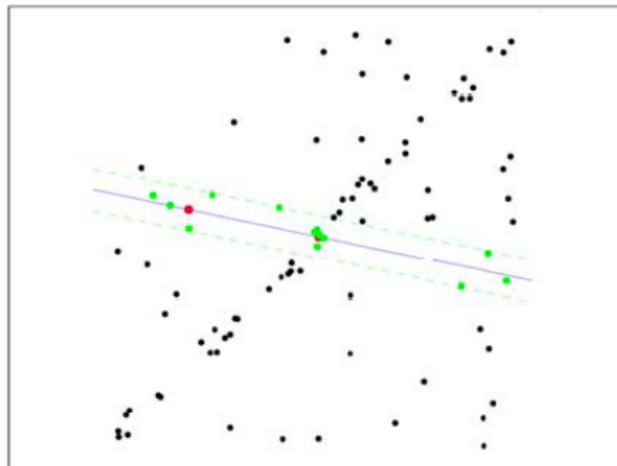
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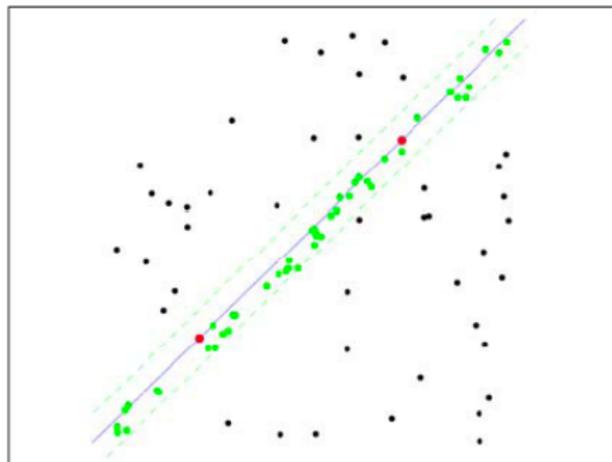
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[Source: R. Raguram]

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# How big is the number of samples?

- For alignment, depends on the motion model
- Each sample is a correspondence (pair of matching points)

Transformation	Matrix	# DoF	Preserves	Icon
translation	$\begin{bmatrix} \mathbf{I} &   & \mathbf{t} \end{bmatrix}_{2 \times 3}$	2	orientation	
rigid (Euclidean)	$\begin{bmatrix} \mathbf{R} &   & \mathbf{t} \end{bmatrix}_{2 \times 3}$	3	lengths	
similarity	$\begin{bmatrix} s\mathbf{R} &   & \mathbf{t} \end{bmatrix}_{2 \times 3}$	4	angles	
affine	$\begin{bmatrix} \mathbf{A} \end{bmatrix}_{2 \times 3}$	6	parallelism	
projective	$\begin{bmatrix} \tilde{\mathbf{H}} \end{bmatrix}_{3 \times 3}$	8	straight lines	

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Next class ... more on cameras and projection