UiO Separtment of Technology Systems

University of Oslo

Estimating homographies from feature correspondences

Thomas Opsahl

2023





Homographies and perspective imaging F_i F_i

The perspective imaging of a planar surface corresponds (exactly!) to a homography

$$\mathbf{H}_{is}\tilde{\mathbf{x}}^{s} = \tilde{\mathbf{u}}^{i}$$

$$\begin{bmatrix} h_{1} & h_{2} & h_{3} \\ h_{4} & h_{5} & h_{6} \\ h_{7} & h_{8} & h_{9} \end{bmatrix} \begin{bmatrix} x \\ y \\ 1 \end{bmatrix} = \begin{bmatrix} u \\ v \\ 1 \end{bmatrix}$$

$$\mathbf{TEK5030}$$



Two overlapping perspective images of a planar scene are "perfectly" related by a homography

Two overlapping perspective images of a planar scene are "perfectly" related by a homography

Two overlapping images from a camera rotating about its projective center (pinhole) are also "perfectly" related by a homography





Hence it is often reasonable to assume that two overlapping images are related by a homography

- Planar or almost planar scene, i.e. when the distance to the scene is relatively much larger than the 3D structures in the scene
- Purely rotating camera or when the distance to the scene is relatively much larger than the camera translation between images







Knowing the homography between two images allows us to map one image into the other image's coordinate system

 H_{ba} : Img_a \rightarrow Img_b

 $H_{ab} = H_{ba}^{-1}$: Img_b \rightarrow Img_a

Coregistering images like this allows us to compare and combine information in overlapping images

Coregistration is therefore an important first step for applications like change detection and image mosaicing







Homography estimation Understanding the problem

We know that

• A homography is a projective transformation that we can represent by a homogeneous 3x3 matrix

 $\mathbf{H}_{ba}\widetilde{\mathbf{u}}^a = \widetilde{\mathbf{u}}^b$

- $\begin{bmatrix} h_1 & h_2 & h_3 \\ h_4 & h_5 & h_6 \\ h_7 & h_8 & h_9 \end{bmatrix} \begin{bmatrix} u^a \\ v^a \\ 1 \end{bmatrix} = \begin{bmatrix} u^b \\ v^b \\ 1 \end{bmatrix}$
- Point correspondences u^a_i ↔ u^b_i can be established automatically, but some wrong correspondences are to be expected







Homography estimation Understanding the problem



This equality involves a homogeneous matrix and two homogenous points/vectors

Since the points are known, we can fix their numerical representation to be the standard representation (last coordinate equal to 1)

The scale ambiguity of the matrix, is however something that we have to take into account when we try to solve for its unknown elements



Homography estimation Understanding the problem



Direct linear transform

The direct linear transformation (DLT) is an algorithm for solving a set of variables from a set of equalities like this (equal up to scale)

By rewriting the equality into a set of proper linear homogeneous equations, we represent the problem in a way that is naturally scale ambiguous

This will allow us to determine the variables using standard methods from linear algebra

$$\begin{bmatrix} h_1 & h_2 & h_3 \\ h_4 & h_5 & h_6 \\ h_7 & h_8 & h_9 \end{bmatrix} \begin{bmatrix} u^a \\ v^a \\ 1 \end{bmatrix} = \begin{bmatrix} u^b \\ v^b \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} h_1 u^a + h_2 v^a + h_3 \\ h_4 u^a + h_5 v^a + h_6 \\ h_7 u^a + h_8 v^a + h_9 \end{bmatrix} = \begin{bmatrix} u^b \\ v^b \\ 1 \end{bmatrix}$$

This equality corresponds to a system of three equations

$$\begin{bmatrix} h_1 & h_2 & h_3 \\ h_4 & h_5 & h_6 \\ h_7 & h_8 & h_9 \end{bmatrix} \begin{bmatrix} u^a \\ v^a \\ 1 \end{bmatrix} = \begin{bmatrix} u^b \\ v^b \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} I \\ II \\ III \\ III \end{bmatrix} \begin{cases} h_1 u^a + h_2 v^a + h_3 = u^b \\ h_4 u^a + h_5 v^a + h_6 = v^b \\ h_7 u^a + h_8 v^a + h_9 = 1 \end{cases}$$

This equality corresponds to a system of three equations

$$\begin{bmatrix} h_1 & h_2 & h_3 \\ h_4 & h_5 & h_6 \\ h_7 & h_8 & h_9 \end{bmatrix} \begin{bmatrix} u^a \\ v^a \\ 1 \end{bmatrix} = \begin{bmatrix} u^b \\ v^b \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} I \\ II \\ III \\ III \end{bmatrix} \begin{cases} h_1 u^a + h_2 v^a + h_3 = u^b \\ h_4 u^a + h_5 v^a + h_6 = v^b \\ h_7 u^a + h_8 v^a + h_9 = 1 \end{cases}$$

We can reformulate this into three linear homogeneous equations

$$\begin{bmatrix} v^{b}I = u^{b}II \\ I = u^{b}III \\ II = v^{b}III \\ II = v^{b}III \end{bmatrix} \begin{cases} h_{1}u^{a}v^{b} + h_{2}v^{a}v^{b} + h_{3}v^{b} = h_{4}u^{a}u^{b} + h_{5}v^{a}u^{b} + h_{6}u^{b} \\ h_{1}u^{a} + h_{2}v^{a} + h_{3} = h_{7}u^{a}u^{b} + h_{8}v^{a}u^{b} + h_{9}u^{b} \\ h_{4}u^{a} + h_{5}v^{a} + h_{6} = h_{7}u^{a}v^{b} + h_{8}v^{a}v^{b} + h_{9}v^{b} \end{cases}$$

$$\begin{bmatrix} h_1 & h_2 & h_3 \\ h_4 & h_5 & h_6 \\ h_7 & h_8 & h_9 \end{bmatrix} \begin{bmatrix} u^a \\ v^a \\ 1 \end{bmatrix} = \begin{bmatrix} u^b \\ v^b \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} I \\ II \\ III \\ III \end{bmatrix} \begin{cases} h_1 u^a + h_2 v^a + h_3 = u^b \\ h_4 u^a + h_5 v^a + h_6 = v^b \\ h_7 u^a + h_8 v^a + h_9 = 1 \end{cases}$$

We can reformulate this into three linear homogeneous equations

$$\begin{cases} u^{a}v^{b}h_{1} + v^{a}v^{b}h_{2} + v^{b}h_{3} - u^{a}u^{b}h_{4} - v^{a}u^{b}h_{5} - u^{b}h_{6} = 0 \\ u^{a}h_{1} + v^{a}h_{2} + 1h_{3} - u^{a}u^{b}h_{7} - v^{a}u^{b}h_{8} - u^{b}h_{9} = 0 \\ u^{a}h_{4} + v^{a}h_{5} + 1h_{6} - u^{a}v^{b}h_{7} - v^{a}v^{b}h_{8} - v^{b}h_{9} = 0 \end{cases}$$



$$\begin{bmatrix} h_1 & h_2 & h_3 \\ h_4 & h_5 & h_6 \\ h_7 & h_8 & h_9 \end{bmatrix} \begin{bmatrix} u^a \\ v^a \\ 1 \end{bmatrix} = \begin{bmatrix} u^b \\ v^b \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} I \\ II \\ III \\ III \end{bmatrix} \begin{cases} h_1 u^a + h_2 v^a + h_3 = u^b \\ h_4 u^a + h_5 v^a + h_6 = v^b \\ h_7 u^a + h_8 v^a + h_9 = 1 \end{cases}$$

Rewrite to get a homogeneous matrix equation

$$\begin{aligned} \left(u^{a}v^{b}h_{1} + v^{a}v^{b}h_{2} + v^{b}h_{3} - u^{a}u^{b}h_{4} - v^{a}u^{b}h_{5} - u^{b}h_{6} + 0h_{7} + 0h_{8} + 0h_{9} = 0 \\ u^{a}h_{1} + v^{a}h_{2} + 1h_{3} + 0h_{4} + 0h_{5} + 0h_{6} - u^{a}u^{b}h_{7} - v^{a}u^{b}h_{8} - u^{b}h_{9} = 0 \\ 0h_{1} + 0h_{2} + 0h_{3} + u^{a}h_{4} + v^{a}h_{5} + 1h_{6} - u^{a}v^{b}h_{7} - v^{a}v^{b}h_{8} - v^{b}h_{9} = 0 \end{aligned}$$

$$\begin{bmatrix} h_{1} & h_{2} & h_{3} \\ h_{4} & h_{5} & h_{6} \\ h_{7} & h_{8} & h_{9} \end{bmatrix} \begin{bmatrix} u^{a} \\ v^{a} \\ 1 \end{bmatrix} = \begin{bmatrix} u^{b} \\ v^{b} \\ 1 \end{bmatrix}$$

$$\begin{pmatrix} u^{a}v^{b} & v^{a}v^{b} & v^{b} & -u^{a}u^{b} & -v^{a}u^{b} & -u^{b} & 0 & 0 \\ u^{a} & v^{a} & 1 & 0 & 0 & 0 & -u^{a}u^{b} & -v^{a}u^{b} & -v^{a}u^{b} & -u^{b} \\ 0 & 0 & 0 & u^{a} & v^{a} & 1 & -u^{a}v^{b} & -v^{a}v^{b} & -v^{b} \end{bmatrix} \begin{bmatrix} h_{1} \\ h_{2} \\ h_{3} \\ h_{4} \\ h_{5} \\ h_{6} \\ h_{7} \\ h_{8} \\ h_{9} \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \end{bmatrix}$$

$$\begin{bmatrix} h_1 & h_2 & h_3 \\ h_4 & h_5 & h_6 \\ h_7 & h_8 & h_9 \end{bmatrix} \begin{bmatrix} u^a \\ v^a \\ 1 \end{bmatrix} = \begin{bmatrix} u^b \\ v^b \\ 1 \end{bmatrix}$$
$$\begin{bmatrix} u^a v^b & v^a v^b & v^b & -u^a u^b & -v^a u^b & -u^b & 0 & 0 & 0 \\ u^a & v^a & 1 & 0 & 0 & 0 & -u^a u^b & -v^a u^b & -u^b \\ 0 & 0 & 0 & u^a & v^a & 1 & -u^a v^b & -v^a v^b & -v^b \end{bmatrix} \mathbf{h} = \mathbf{0}$$

The nature of homogeneous equations implies that all solutions are scale ambiguous, i.e. if $Ah^* = 0$, then we also have that $A(\alpha h^*) = 0$ for any non-zero scalar α

This means that we can use standard methods in linear algebra to determine h

$$\begin{bmatrix} h_{1} & h_{2} & h_{3} \\ h_{4} & h_{5} & h_{6} \\ h_{7} & h_{8} & h_{9} \end{bmatrix} \begin{bmatrix} u^{a} \\ v^{a} \\ 1 \end{bmatrix} = \begin{bmatrix} u^{b} \\ v^{b} \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} u^{a}v^{b} & v^{a}v^{b} & v^{b} & -u^{a}u^{b} & -v^{a}u^{b} & -u^{b} & 0 & 0 & 0 \\ u^{a} & v^{a} & 1 & 0 & 0 & 0 & -u^{a}u^{b} & -v^{a}u^{b} & -u^{b} \\ 0 & 0 & 0 & u^{a} & v^{a} & 1 & -u^{a}v^{b} & -v^{a}v^{b} & -v^{b} \end{bmatrix} \mathbf{h} = \mathbf{0}$$

At first glance, it seems like each point correspondence $\mathbf{u}_i^a \leftrightarrow \mathbf{u}_i^b$ puts three constraints on **h**

$$\begin{bmatrix} h_{1} & h_{2} & h_{3} \\ h_{4} & h_{5} & h_{6} \\ h_{7} & h_{8} & h_{9} \end{bmatrix} \begin{bmatrix} u^{a} \\ v^{a} \\ 1 \end{bmatrix} = \begin{bmatrix} u^{b} \\ v^{b} \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} u^{a}v^{b} & v^{a}v^{b} & v^{b} & -u^{a}u^{b} & -v^{a}u^{b} & -u^{b} & 0 & 0 & 0 \\ u^{a} & v^{a} & 1 & 0 & 0 & 0 & -u^{a}u^{b} & -v^{a}u^{b} & -u^{b} \\ 0 & 0 & 0 & u^{a} & v^{a} & 1 & -u^{a}v^{b} & -v^{a}v^{b} & -v^{b} \end{bmatrix} \mathbf{h} = \mathbf{0}$$

At first glance, it seems like each point correspondence $\mathbf{u}_i^a \leftrightarrow \mathbf{u}_i^b$ puts three constraints on **h**

They are however not independent: $v^b \cdot row_2 - u^b \cdot row_3 = row_1$

$$\begin{bmatrix} h_{1} & h_{2} & h_{3} \\ h_{4} & h_{5} & h_{6} \\ h_{7} & h_{8} & h_{9} \end{bmatrix} \begin{bmatrix} u^{a} \\ v^{a} \\ 1 \end{bmatrix} = \begin{bmatrix} u^{b} \\ v^{b} \\ 1 \end{bmatrix}$$

$$\begin{pmatrix} u^{a}v^{b} & v^{a}v^{b} & v^{b} & -u^{a}u^{b} & -v^{a}u^{b} & -u^{b} & 0 & 0 \\ u^{a} & v^{a} & 1 & 0 & 0 & 0 & -u^{a}u^{b} & -v^{a}u^{b} & -u^{b} \\ 0 & 0 & 0 & u^{a} & v^{a} & 1 & -u^{a}v^{b} & -v^{a}v^{b} & -v^{b} \end{bmatrix} \mathbf{h} = \mathbf{0}$$

At first glance, it seems like each point correspondence $\mathbf{u}_i^a \leftrightarrow \mathbf{u}_i^b$ puts three constraints on **h**

They are however not independent: $v^b \cdot row_2 - u^b \cdot row_3 = row_1$

Each point correspondence $\mathbf{u}_i^a \leftrightarrow \mathbf{u}_i^b$ corresponds to two constraints on **h**, so we can choose to disregard one of them (does not matter which one)

$$\begin{bmatrix} u_1^a & v_1^a & 1 & 0 & 0 & 0 & -u_1^a u_1^b & -v_1^a u_1^b & -u_1^b \\ 0 & 0 & 0 & u_1^a & v_1^a & 1 & -u_1^a v_1^b & -v_1^a v_1^b & -v_1^b \\ u_2^a & v_2^a & 1 & 0 & 0 & 0 & -u_2^a u_2^b & -v_2^a u_2^b & -u_2^b \\ 0 & 0 & 0 & u_2^a & v_2^a & 1 & -u_2^a v_2^b & -v_2^a v_2^b & -v_2^b \\ \vdots & \vdots \\ u_n^a & v_n^a & 1 & 0 & 0 & 0 & -u_n^a u_n^b & -v_n^a u_n^b & -u_n^b \\ 0 & 0 & 0 & u_n^a & v_n^a & 1 & -u_n^a v_n^b & -v_n^a v_n^b & -v_n^b \end{bmatrix} \mathbf{h} = \mathbf{0}$$

In a situation with several point correspondences, $\{\mathbf{u}_i^a \leftrightarrow \mathbf{u}_i^b\}_{i=1,...,n}$, we can combine all of them into a $2n \times 9$ matrix **A**

Since we are not after the trivial solution, h = 0, it is clear that h must lie in the null space of A

There are several ways to determine the null space of a matrix, but we will use singular value decomposition (SVD)

Singular Value Decomposition (SVD)

SVD is a factorization of a $m \times n$ matrix **A** into a product $\mathbf{A} = \mathbf{USV}^T$

Where

U is a $m \times m$ orthogonal matrix **S** is a m $\times n$ rectangular diagonal matrix **V** is a $n \times n$ orthogonal matrix

The non-zero diagonal entries of **S** are known as the singular values of **A**

The columns of **U** and **V** are known as leftsingular vectors and right-singular vectors correspondingly





Singular Value Decomposition (SVD)

SVD is a factorization of a $m \times n$ matrix **A** into a product $\mathbf{A} = \mathbf{USV}^T$

Where

U is a $m \times m$ orthogonal matrix **S** is a m × 9 rectangular diagonal matrix **V** is a 9 × 9 orthogonal matrix

The non-zero diagonal entries of **S** are known as the singular values of **A**

The columns of **U** and **V** are known as leftsingular vectors and right-singular vectors correspondingly For homography estimation: n = 9 and $m \ge 8$

$$\mathbf{S} = \begin{bmatrix} s_1 & & \mathbf{0} \\ & \ddots & \mathbf{0} \\ & \mathbf{0} & & s_9 \end{bmatrix}$$

 $\mathbf{U} = [\mathbf{u}_1, \mathbf{u}_2, \dots, \mathbf{u}_m]$

 $\mathbf{V} = [\mathbf{v}_1, \mathbf{v}_2, \dots, \mathbf{v}_9]$



Singular Value Decomposition (SVD)

The factorization $\mathbf{A} = \mathbf{U}\mathbf{S}\mathbf{V}^T$ is not unique, but one can always choose a factorization such that the diagonal entries of \mathbf{S} are in descending order

Key result

The right singular vectors corresponding to vanishing singular values, i.e. $s_i = 0$, spans the null space of **A**

This means that, if **A** has rank 8 and the diagonal entries in **S** are in descending order, we have that $s_9 = 0$ and so $\mathbf{h} = \mathbf{v}_9$

For homography estimation: n = 9 and $m \ge 8$



 $\mathbf{U} = [\mathbf{u}_1, \mathbf{u}_2, \dots, \mathbf{u}_m]$

 $\mathbf{V} = [\mathbf{v}_1, \mathbf{v}_2, \dots, \mathbf{v}_9]$

Since each point correspondence $\mathbf{u}_i^a \leftrightarrow \mathbf{u}_i^b$ provides 2 rows in matrix **A**, we need at least 4 correspondences for **A** to have rank 8

Given 4 point correspondences, we are guaranteed **A** to have rank 8 as long as no three points (in either of the two images) are co-linear



In principle, the matrix A should have rank 8 even when we have more than 4 point correspondences

But, errors in the feature matching and uncertainties in the actual positioning of the feature points will typically cause **A** to have rank 9



Analyzing matrix A's singular values should however reveal that the 9th and smallest singular value is close to 0 while the other 8 singular values are not

This indicates that $\mathbf{h} = \mathbf{v}_9$ still will be a reasonable solution to the problem





Key result

The right-singular vector corresponding to the smallest singular value, is the optimal solution to Ah = 0 in the sense that it minimizes the total least squares ||Ah|| under the constraint ||h|| = 1





When estimating the homography from only 4 point correspondences in a non-degenerate configuration, A will always have rank 8 and its smallest singular value will always be 0

So by choosing $h = v_9$ we are guaranteed that $||Ah|| \equiv 0$ (at least up to numerical precision)

This means that we are guaranteed to find a perfect homography for the 4 given point correspondences!



- 2. Obtain the SVD of $A: A = USV^T$
- 3. If **S** is diagonal with positive values in descending order along the main diagonal, then \mathbf{h} equals the last column of \mathbf{V}
- 4. Reconstruct \mathbf{H}_{ba} from \mathbf{h}

$$\mathbf{H}_{ba} = \begin{bmatrix} h_1 & h_2 & h_3 \\ h_4 & h_5 & h_6 \\ h_7 & h_8 & h_9 \end{bmatrix}$$

For more than 4 point correspondences, the basic DLT algorithm is however not recommended

The terms of matrix **A** will in general be of very different orders of magnitude $[10^0, 10^6]$ This causes errors in the point positions to have very different impact on the estimation

To alleviate this, it is custom perform the estimation on normalized image coordinates instead



For more than 4 point correspondences, the basic DLT algorithm is however not recommended

The terms of matrix **A** will in general be of very different orders of magnitude $[10^0, 10^6]$ This causes errors in the point positions to have very different impact on the estimation

To alleviate this, it is custom perform the estimation on normalized image coordinates instead

The normalizing transformations can be created in several ways, e.g. based on the mean vector μ and covariance matrix Σ of the two point sets { (u_i^a, v_i^a) } and { (u_i^b, v_i^b) }

$$\mathbf{T}_{a_n a} = \begin{bmatrix} \mathbf{\Sigma}_a^{-\frac{1}{2}} & -\mathbf{\Sigma}_a^{-\frac{1}{2}} \boldsymbol{\mu}_a \\ \mathbf{0} & 1 \end{bmatrix} \qquad \qquad \mathbf{T}_{b_n b} = \begin{bmatrix} \mathbf{\Sigma}_b^{-\frac{1}{2}} & -\mathbf{\Sigma}_b^{-\frac{1}{2}} \boldsymbol{\mu}_b \\ \mathbf{0} & 1 \end{bmatrix}$$



For more than 4 point correspondences, the basic DLT algorithm is however not recommended

The terms of matrix **A** will in general be of very different orders of magnitude $[10^0, 10^6]$ This causes errors in the point positions to have very different impact on the estimation

To alleviate this, it is custom perform the estimation on normalized image coordinates instead

The normalizing transformations can be created in several ways, e.g. based on the mean vector μ and covariance matrix Σ of the two point sets { (u_i^a, v_i^a) } and { (u_i^b, v_i^b) }

$$\mathbf{T}_{a_n a} = \begin{bmatrix} \mathbf{\Sigma}_a^{-\frac{1}{2}} & -\mathbf{\Sigma}_a^{-\frac{1}{2}} \mathbf{\mu}_a \\ \mathbf{0} & \mathbf{1} \end{bmatrix} \qquad \mathbf{M} \text{a trix square root of } \mathbf{\Sigma}^{-1} \qquad \mathbf{J} \qquad \mathbf{$$





How do we know if an estimated homography is good or bad?

We really want to estimate the homography in a RANSAC scheme, but how can we determine if a given point correspondence $(u_i^a, v_i^a) \leftrightarrow (u_i^b, v_i^b)$ is an inlier or an outlier?



How do we know if an estimated homography is good or bad?

We really want to estimate the homography in a RANSAC scheme, but how can we determine if a given point correspondence $(u_i^a, v_i^a) \leftrightarrow (u_i^b, v_i^b)$ is an inlier or an outlier?

Algebraic error for a point correspondence $(u_i^a, v_i^a) \leftrightarrow (u_i^b, v_i^b)$

$$\varepsilon_{i} = \|\mathbf{A}_{i}\mathbf{h}\| \text{ where } \mathbf{A}_{i} = \begin{bmatrix} u_{i}^{a} & v_{i}^{a} & 1 & 0 & 0 & 0 & -u_{i}^{a}u_{i}^{b} & -v_{i}^{a}u_{i}^{b} & -u_{i}^{b} \\ 0 & 0 & 0 & u_{i}^{a} & v_{i}^{a} & 1 & -u_{i}^{a}v_{i}^{b} & -v_{i}^{a}v_{i}^{b} & -v_{i}^{b} \end{bmatrix}$$

Total squared algebraic error for the homography

$$\varepsilon^2 = \sum_i \|\mathbf{A}_i \mathbf{h}\|^2 = \|\mathbf{A}\mathbf{h}\|^2$$

Geometric error for a point correspondence $(u_i^a, v_i^a) \leftrightarrow (u_i^b, v_i^b)$

 $\varepsilon_i = d\left(\mathbf{u}_i^a, \mathbf{H}_{ba}^{-1}\left(\mathbf{u}_i^b\right)\right) + d\left(\mathbf{u}_i^b, \mathbf{H}_{ba}\left(\mathbf{u}_i^a\right)\right)$ where $d(\cdot, \cdot)$ is the Euclidean distance

Total squared geometric error for the homography

$$\varepsilon^2 = \sum_i \varepsilon_i^2$$

Geometric error for a point correspondence $(u_i^a, v_i^a) \leftrightarrow (u_i^b, v_i^b)$

 $\varepsilon_i = d\left(\mathbf{u}_i^a, \mathbf{H}_{ba}^{-1}\left(\mathbf{u}_i^b\right)\right) + d\left(\mathbf{u}_i^b, \mathbf{H}_{ba}\left(\mathbf{u}_i^a\right)\right)$ where $d(\cdot, \cdot)$ is the Euclidean distance

Total squared geometric error for the homogra If we're only concerned with the error in one of the images,

$$\varepsilon^2 = \sum_i \varepsilon_i^2$$

we can also consider a one-sided geometric error

$$\varepsilon_i = d\left(\mathbf{u}_i^a, \mathbf{H}_{ba}^{-1}(\mathbf{u}_i^b)\right)$$

 $\varepsilon_i = d\left(\mathbf{u}_i^b, \mathbf{H}_{ba}(\mathbf{u}_i^a)\right)$



Algebraic error

- $\varepsilon_i = \|\mathbf{A}_i \mathbf{h}\|$
- Not physically meaningful
- Estimating the homography with minimal algebraic error, is easy (DLT) and well suited for use in a RANSAC estimation scheme

Geometric error

- $\varepsilon_i = d\left(\mathbf{u}_i^a, \mathbf{H}_{ba}^{-1}\left(\mathbf{u}_i^b\right)\right) + d\left(\mathbf{u}_i^b, \mathbf{H}_{ba}\left(\mathbf{u}_i^a\right)\right)$
- Physically meaningful
- Estimating the homography with minimal geometric error is a non-linear least squares problem and requires iterative estimation techniques



Homography estimation RANSAC

Algorithm – RANSAC

For a set of point-correspondences $S = \{(u_i^a, v_i^a) \leftrightarrow (u_i^b, v_i^b)\}$, perform N iterations

- 1. Compute $H_{ba,tst}$ for 4 random correspondences with the DLT algorithm
- 2. Determine the set of inliers for $H_{ba,tst}$

$$S_{tst} = \left\{ (u_i^a, v_i^a) \leftrightarrow \left(u_i^b, v_i^b \right) s. t. \ \varepsilon_i < t \right\}$$

Here ε_i is the geometric error and t is a chosen value for max acceptable error

3. If S_{tst} is the largest set of inliers so far

$$\begin{split} S_{IN} &= S_{tst} \\ \mathbf{H}_{ba} &= \mathbf{H}_{ba,tst} \\ N &= \frac{\log(1-p)}{\log(1-\omega^n)} \quad \text{where } \omega = \frac{|S_{IN}|}{|S|}, n = 4 \text{ and } p = 0.99 \end{split}$$

Afterwards, estimate the homography H_{ba} based on only inlier-correspondences

- Minimal algebraic error normalized DLT
- Minimal geometric error iterative, non-linear least squares



Let us compose these two images into a larger image, an image mosaic





We start by finding key points and representing them by descriptors





Establish point-correspondences by matching descriptors





Some bad matches!

Establish point-correspondences by matching descriptors







Determine the inlier set of point correspondences by estimating the homography in a RANSAC scheme

H_{ba}

Estimate the homography H_{ba} based on inlier correspondences only





Now we could use the homography H_{ba} to transform (warp) the left image into the right image's coordinate system

 \mathbf{H}_{ba}

But, if we want to see the full mosaic it is necessary to transform both images into a more suitable coordinate system



Here we've chosen to transform both images into a shifted version of the right image's coordinate system

If \mathbf{T}_{mb} is the transformation from the right image to this new "mosaic image", the transformation of the left image is given by $\mathbf{T}_{mb}\mathbf{H}_{ba}$



Once both images are represented in the same coordinates, we can choose to compose them in several different ways







Blending with a ramp + histogram equalization

Summary

Often reasonable to assume that two overlapping perspective images are related by a homography

- Planar or almost planar scene
- Purely rotating or almost purely rotating camera

Homography estimation

- 1. Establish point correspondences
- 2. RANSAC estimation of homography to remove bad correspondences
- 3. Estimate homography based on good correspondences
 - Minimal algebraic error: Normalized DLT
 - Minimal geometric error: Iterative, non-linear least squares

Image mosaic from two images

The homography can be used to transform both images into a common coordinate frame











Supplementary material

Recommended

- Richard Szeliski: Computer Vision: Algorithms and Applications 2nd ed
 - Chapter 8 "Image alignment and stitching" and in particular section 8.2 about "image stitching"

