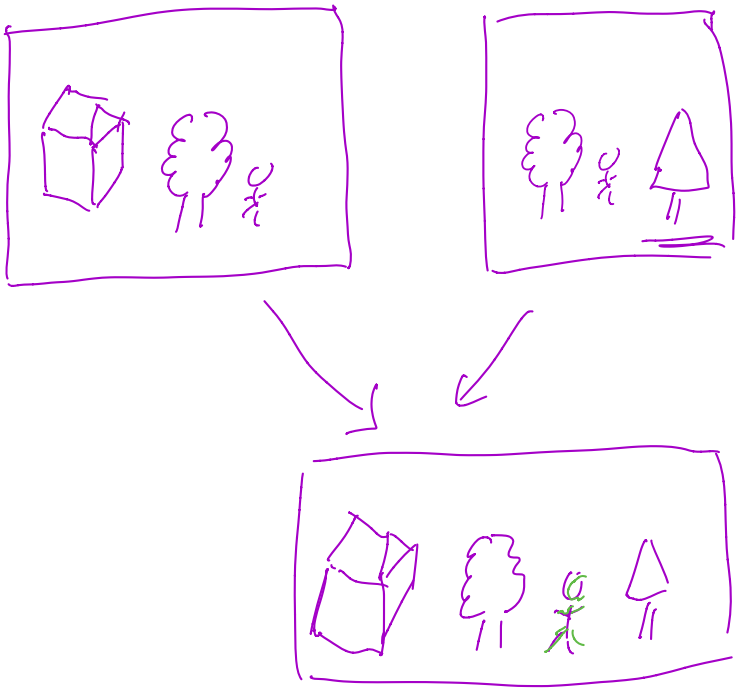
HW 4 posted, due Tue 3/23
Lec 11/12 LE problem out by 5:30 EST due Wed night

CSci 4270 and 6270
Computational Vision, SPring 2021
Lecture 12: Two Image Matching
March 8, 2021

Building a Montage or Mosaic from Two (or More) Images

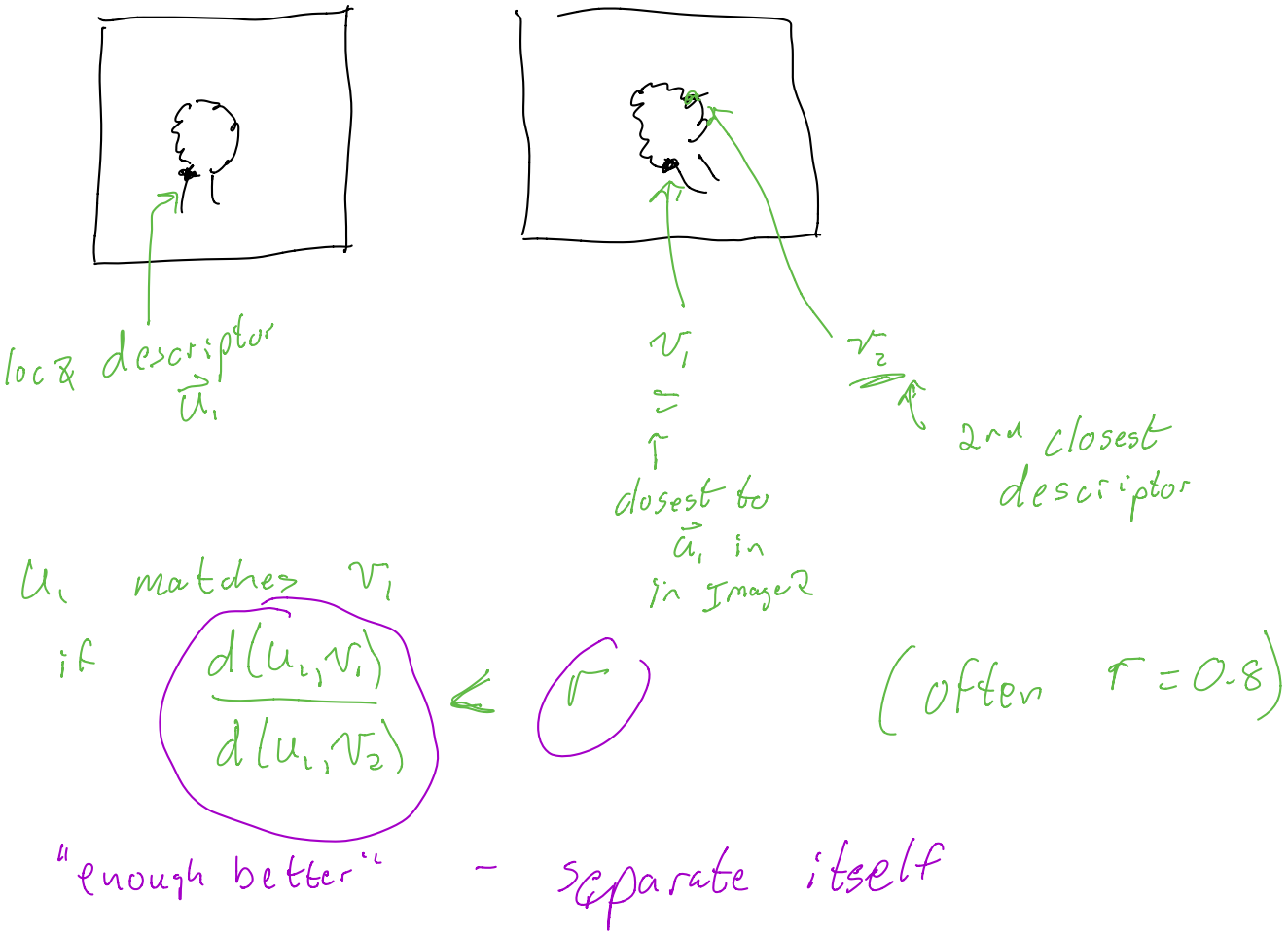
We will investigate the problem of estimating inter-image transformations in the context of matching two (or more) images and forming a mosaic. Here are the steps:

1. Keypoint extraction and description in each image ✓
2. Keypoint matching between images ✓
3. Using the best matches ✓
4. Estimating the transformation parameters between two images.
5. Eliminating false matches
6. Estimating the transformation parameters for all images.
7. Mapping the images and blending



Extracting and Matching Keypoints

- Extract keypoints and descriptors in each image.
- For each keypoint, find the keypoints with the two best matching descriptors in the other image and compute the ratio score.
- Keep the "best" matches — those with SIFT descriptor ratio scores below 0.8.
- Not every keypoint is used — in fact most are not
- Even among those with ratio below 0.8, some of the matches can be wrong, having a disastrous effect on the transformation.
- Before we consider how to handle these, we will focus on the least-squares estimation assuming the correspondences are correct.

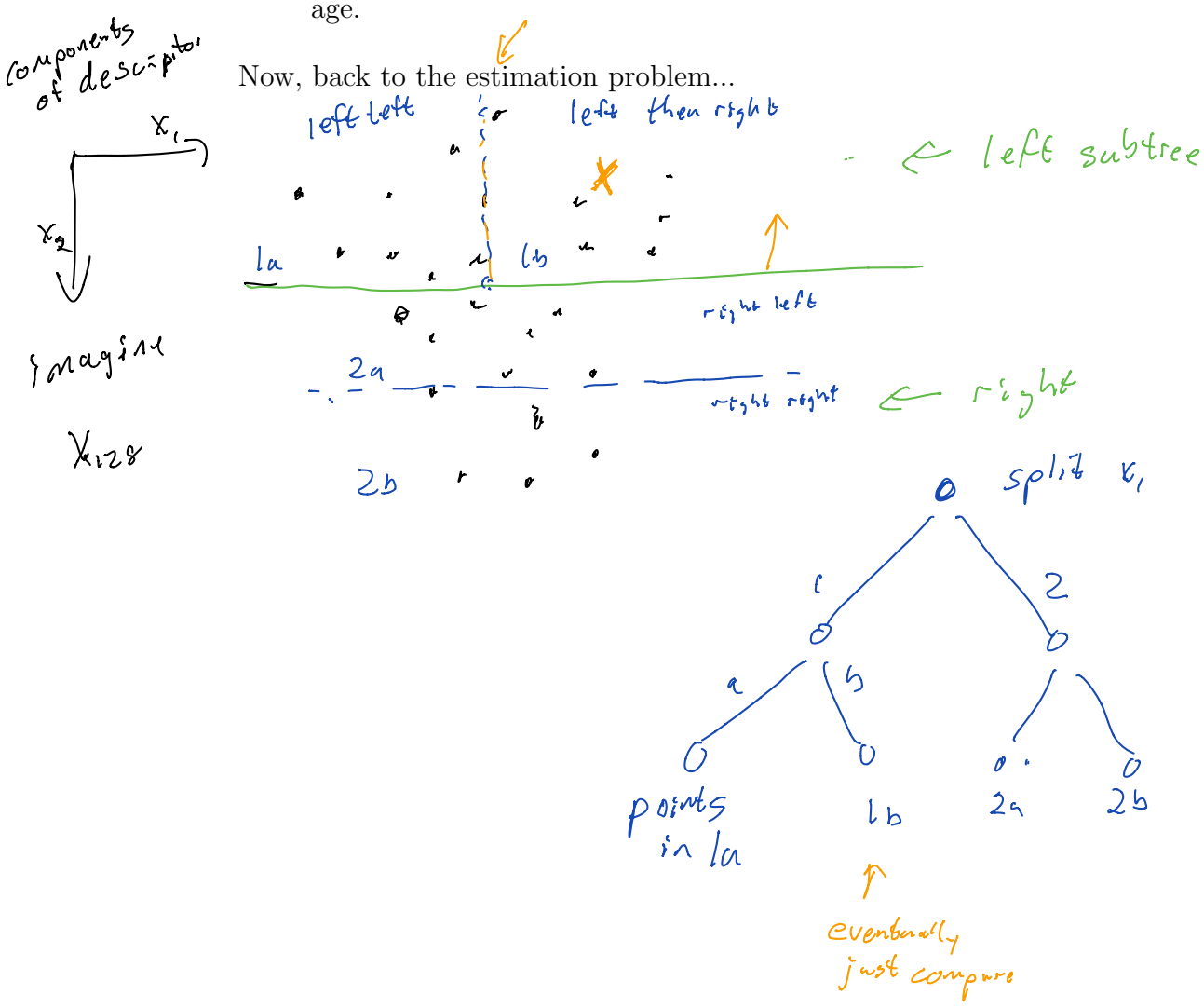


Aside: K-D Tree

- With 2,000 keypoints or more keypoints in each, the quadratic all-against-all matching process is expensive.
- Need a data structure to make searching fast.
- Main choice is a K-D tree, a multidimensional version of a binary search tree

- General idea:
  - At each node of the tree, split the list of keypoint descriptor vectors to be stored in half based on the median value of one of the vector dimensions (128 in our case).
  - Continue recursively
- Recursion stops at nodes where the number of vectors to stored is lower than a threshold (e.g. about 5-10)
- Searching becomes a prioritized exploration of the nodes of the tree.
- Search time is practically about $O(\sqrt{N})$.
- Implementation of this and alternative versions are in the FLANN package.

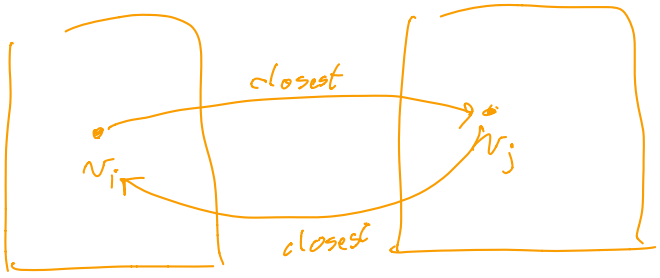
Now, back to the estimation problem...



ORB Descriptor

Rublee, Rabaud, Konolige and Bradski, *ORB: An efficent alternative to SIF or SURF*, ICCV 2011

- One of many alternatives to SIFT.
- 32 bytes per descriptor; SIFT is 128 (or more)
- Distance measure between two descriptors is the Hamming distance — the number of bit differences across as $32 \times 8 = 256$ bit
  - Contrast this to the Euclidean distance metric used for SIFT keypoints.
- The SIFT ratio test is often replaced by a *symmetic matching test*:
  - Let $\mathbf{v}_i$ be the ORB descriptor for the i-th keypoint from image I_1.
  - Let $\mathbf{v}_j$ be the ORB descriptor for the j-th keypoint from image I_2.
  - Keypoints i and j are labeled as a match if $\mathbf{v}_j$ is the closest descriptor from I_2 to $\mathbf{v}_i$ **and** if $\mathbf{v}_i$ is the closest descriptor to $\mathbf{v}_j$ from image I_2.
- For HW 4 we recommend using SIFT instead of ORB.



Formulating the Estimation Problem

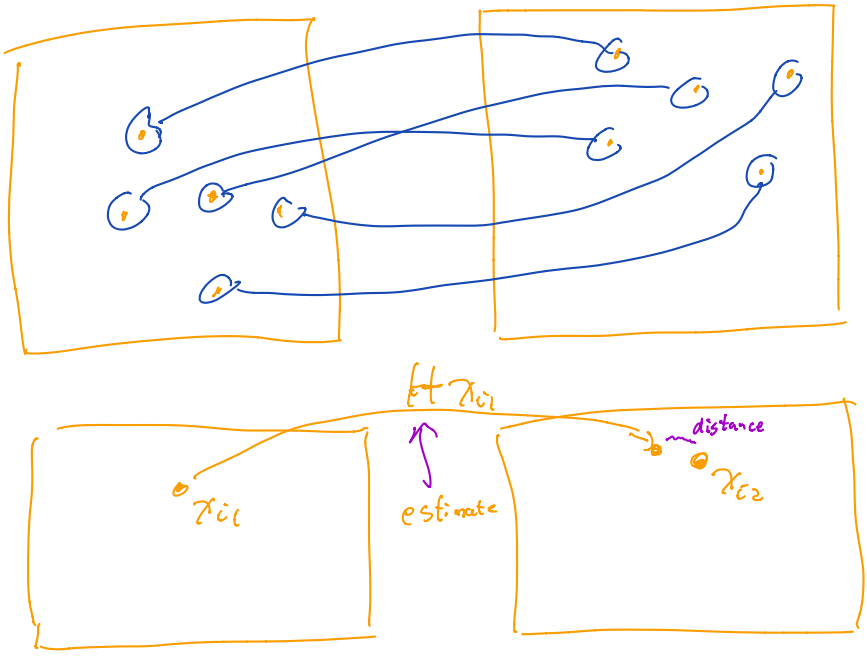
- Given are sets of corresponding keypoint locations between two images, I_1 and I_2, as selected by matching descriptors.
- Shifting notation a bit from what we've just used, the set of N matches is $\{(\tilde{\mathbf{x}}_{i,1}, \tilde{\mathbf{x}}_{i,2})\}$ for $i = 1, \ldots, N$.
  - These are homogeneous (image) coordinate vectors of pixel locations:

$$\tilde{\mathbf{x}}_{i,1} = \begin{pmatrix} x_{i,1} \\ y_{i,1} \\ 1 \end{pmatrix} \qquad \text{and} \qquad \tilde{\mathbf{x}}_{i,2} = \begin{pmatrix} x_{i,2} \\ y_{i,2} \\ 1 \end{pmatrix}$$

- The problem we want to solve is to find the parameters of the 3×3 homography matrix $\mathbf{H}$ that minimizes the sum of the square distances between

$$\mathbf{H}\tilde{\mathbf{x}}_{i,1} \qquad \text{and} \qquad \tilde{\mathbf{x}}_{i,2}.$$

goal is estimate H



Difficulties This Poses

- Measuring the distance for homogeneous coordinates
- Dealing with the resulting non-linearities
- Different meanings of the different parameters
- Restructuring the equations

We'll start with the simpler problem of the affine transfomation

Estimating the Affine Transformation Parameters

In this case, we can easily use image distances

- Measuring the distance in image I_2, we obtain the least-squares objective function:

$$E(\mathbf{a}) = \sum_i \|\mathbf{x}_{i,2} - (\mathbf{A}\mathbf{x}_{i,1} + \mathbf{t})\|^2$$

- The components of $\mathbf{A}$ and $\mathbf{t}$, combined into $\mathbf{a}$, are unknown and must be estimated by minimizing this equation.
- In detail:
  - $\mathbf{x}_{i,2}$ and $\mathbf{x}_{i,1}$ are non-homogeneous keypoint (pixel) locations:

$$\mathbf{x}_{i,1} = \begin{pmatrix} x_{i,1} \\ y_{i,1} \end{pmatrix} \quad \text{and} \quad \mathbf{x}_{i,2} = \begin{pmatrix} x_{i,2} \\ y_{i,2} \end{pmatrix}$$

  - $\mathbf{A}$ is the 2×2 matrix of affine terms:

$$\mathbf{A} = \begin{pmatrix} a_{1,1} & a_{1,2} \\ a_{2,1} & a_{2,2} \end{pmatrix}$$

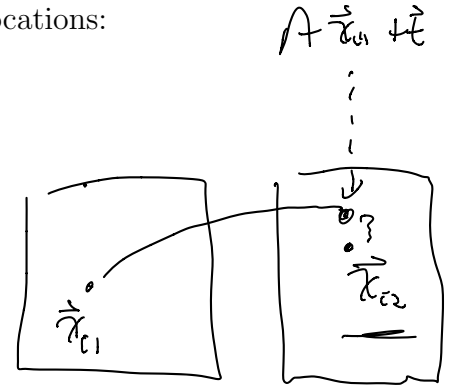
  - $\mathbf{t}$ is the 2×1 vector of translation terms:

$$\mathbf{t} = \begin{pmatrix} t_x \\ t_y \end{pmatrix}$$

  - And finally, $\mathbf{a}$, written as a row vector is

$$\mathbf{a}^\top = \begin{pmatrix} a_{1,1} & a_{1,2} & t_x & a_{2,1} & a_{2,2} & t_y \end{pmatrix}$$

The reasoning for the ordering in $\mathbf{a}$ will be made clear below.



unknowns are a's and t_x t_y

$$\left\| \begin{pmatrix} x_{i2} \\ y_{i2} \end{pmatrix} - \left(\begin{pmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{pmatrix} \begin{pmatrix} x_{i1} \\ y_{i1} \end{pmatrix} + \begin{pmatrix} t_x \\ t_y \end{pmatrix} \right) \right\|^2$$

$$= \left\| \begin{pmatrix} x_{i2} \\ y_{i2} \end{pmatrix} - \begin{pmatrix} a_{11} x_{i1} + a_{12} y_{i1} + t_x \\ a_{21} x_{i1} + a_{22} y_{i1} + t_y \end{pmatrix} \right\|^2$$

$$= \left\| \begin{pmatrix} x_{i2} \\ y_{i2} \end{pmatrix} - \begin{pmatrix} x_{i1} & y_{i1} & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & x_{i1} & y_{i1} & 1 \end{pmatrix} \begin{pmatrix} a_{11} \\ a_{12} \\ t_x \\ a_{21} \\ a_{22} \\ t_y \end{pmatrix} \right\|^2$$

Completing the Estimation

- A little bit of algebra allows us to re-arrange the summation to rewrite the object function as

$$E(\mathbf{a}) = \sum_i \|\mathbf{x}_{i,2} - \mathbf{X}_{i,1}\mathbf{a}\|^2$$

where

$$\mathbf{X}_{i,1} = \begin{pmatrix} x_{i,1} & y_{i,1} & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & x_{i,1} & y_{i,1} & 1 \end{pmatrix}$$

- Computing the derivative of this with respect to the vector $\mathbf{a}$, setting the result equal to $\mathbf{0}$, and solving yields the estimate, $\hat{\mathbf{a}}$:

$$\hat{\mathbf{a}} = (\sum_i \mathbf{X}_{i,1}^\top \mathbf{X}_{i,1})^{-1} (\sum_i \mathbf{X}_{i,1}^\top \mathbf{x}_{i,2})$$

- Of course we solve this using the singular value decomposition (SVD)
- From the components of the estimate $\hat{\mathbf{a}}$ we fill in the values of $\mathbf{A}$ and $\mathbf{t}$.

Discussing the Affine Solution

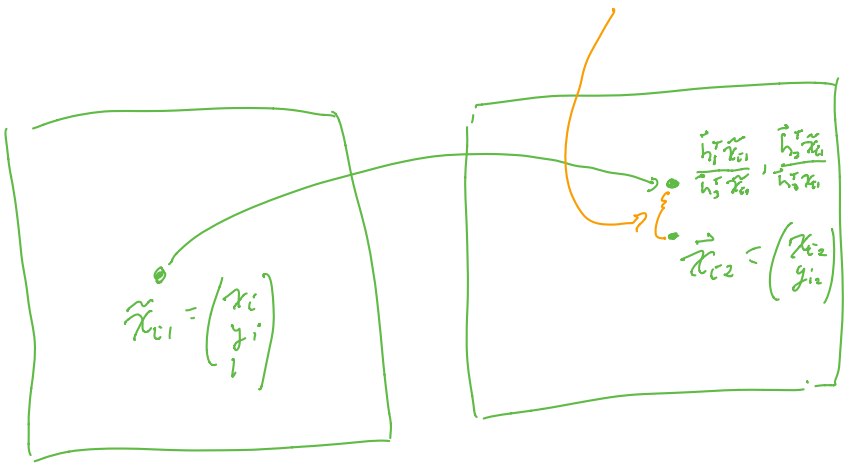
- The derivation (which we will work on in class) illustrates the importance of vector and linear algebra.
- The inversion of the 6×6 matrix $\sum_i \mathbf{X}_{i,1}^\top \mathbf{X}_{i,1}$ may be split into the inversion of two identical 3×3 matrices,
- Whenever inverting matrices to solve estimation problems, we need to be careful of the relative size of the terms of the matrix, as we will discuss in class.

Turning to the Homography Estimation Problem

- Recall that the correspondence set is $\{(\tilde{\mathbf{x}}_{i,1}, \tilde{\mathbf{x}}_{i,2})\}$.
  - Here we are back to homogeneous coordinates.
- We need to estimate the parameters of the 3×3 matrix $\mathbf{H}$ minimizing the distance between the $\tilde{\mathbf{x}}_{i,2}$ image location and the transformed image location $\mathbf{H}\tilde{\mathbf{x}}_{i,1}$.

Measuring the Distance

- Remember, in order to convert $\mathbf{H}\tilde{\mathbf{x}}_{i,1}$ back to affine coordinates, we need to divide by the third row:
  - We write
$$\mathbf{H} = \begin{pmatrix} h_{11} & h_{12} & h_{13} \\ h_{21} & h_{22} & h_{23} \\ h_{31} & h_{32} & h_{33} \end{pmatrix} = \begin{pmatrix} \mathbf{h}_1^\top \\ \mathbf{h}_2^\top \\ \mathbf{h}_3^\top \end{pmatrix}$$
    where $\mathbf{h}_j^\top = \begin{pmatrix} h_{j1} & h_{j2} & h_{j3} \end{pmatrix}$ is row j of $\mathbf{H}$.
  - Now, recall that with homographies, the mapping of a pixel location in image 1 is
$$\begin{pmatrix} u \\ v \\ w \end{pmatrix} = \mathbf{H}\tilde{\mathbf{x}} = \begin{pmatrix} \mathbf{h}_1^\top \tilde{\mathbf{x}} \\ \mathbf{h}_2^\top \tilde{\mathbf{x}} \\ \mathbf{h}_3^\top \tilde{\mathbf{x}} \end{pmatrix}$$
    and the final mapped pixel location is
$$\left(\frac{u}{w}, \frac{v}{w} \right) = \left(\frac{\mathbf{h}_1^\top \tilde{\mathbf{x}}}{\mathbf{h}_3^\top \tilde{\mathbf{x}}}, \frac{\mathbf{h}_2^\top \tilde{\mathbf{x}}}{\mathbf{h}_3^\top \tilde{\mathbf{x}}} \right).$$
- Using this notations, the distance between the mapping of $\tilde{\mathbf{x}}_{i,1}$ into image 2 and the corresponding point location $\tilde{\mathbf{x}}_{i,2}$ in image 2 is
$$d(\tilde{\mathbf{x}}_{i,2}, \mathbf{H}\tilde{\mathbf{x}}_{i,1}) = \left[\left(x_{i,2} - \frac{\mathbf{h}_1^\top \tilde{\mathbf{x}}_{i,1}}{\mathbf{h}_3^\top \tilde{\mathbf{x}}_{i,1}} \right)^2 + \left(y_{i,2} - \frac{\mathbf{h}_2^\top \tilde{\mathbf{x}}_{i,1}}{\mathbf{h}_3^\top \tilde{\mathbf{x}}_{i,1}} \right)^2 \right]^{1/2}$$
- Finally, the objective function is then
$$\sum_i d(\tilde{\mathbf{x}}_{i,2}, \mathbf{H}\tilde{\mathbf{x}}_{i,1})^2$$



Problems Introduced By This Formulation

- Two problems:
  1. Scaling **H** by a non-zero constant has no effect on the distance.
  2. Unknown components of **H** appear in the denominator.
- The first problem is solved by imposing the constraint that the "Frobenius norm" (the sum of the squares of the matrix parameters) is 1.
  - This is likely making a vector into a unit vector.
- The second problem has two different solutions, which we will study in turn:
  1. Clear the denominator and solve the resulting problem. This produces what is called the "algebraic" distance.
  2. Directly solve the non-linear optimization problem.

$$\left\{\left(x_{i2} - \frac{h_1^T \tilde{x}_{i1}}{h_3^T \tilde{x}_{i1}}\right)^2 + \left(y_{i2} - \frac{h_2^T \tilde{x}_{i1}}{h_3^T \tilde{x}_{i1}}\right)^2\right\}^{1/2}$$

equal

$$\left(\frac{1}{h_3^T \tilde{x}_{i1}}\right)\left\{\left(x_{i2} h_3^T \tilde{x}_{i1} - h_1^T \tilde{x}_{i1}\right)^2 + \left(y_{i2} h_3^T \tilde{x}_{i1} - h_2^T \tilde{x}_{i1}\right)^2\right\}^{1/2}$$

make this "1" → essentially ignore this

Forming and Minimizing the Algebraic Distance

- "Clear" the denominator, giving an objective function of

$$\sum_i (x_{i,2}\mathbf{h}_3^\top \tilde{\mathbf{x}}_{i,1} - \mathbf{h}_1^\top \tilde{\mathbf{x}}_{i,1})^2 + (y_{i,2}\mathbf{h}_3^\top \tilde{\mathbf{x}}_{i,1} - \mathbf{h}_2^\top \tilde{\mathbf{x}}_{i,1})^2.$$

$= \sum \|X_i \vec{h}\|^2$ (2×9)

- We now minimize this subject to the Frobenius constraint, which we can write as

$$\mathbf{h}_1^\top \mathbf{h}_1 + \mathbf{h}_2^\top \mathbf{h}_2 + \mathbf{h}_3^\top \mathbf{h}_3 = 1.$$

sum of sq magnitudes are 1

- We will work through the details (a lot) in class.

X_i

$-x_{i1}$	$-y_{i1}$	-1	0	0	0	$x_{i2}x_{i1}$	$x_{i2}y_{i1}$	x_{i2}
0	0	0	$-x_{i1}$	$-y_{i1}$	-1	$y_{i2}x_{i1}$	$y_{i2}y_{i1}$	y_{i2}

$\vec{h}$

h_{11} h_{12} h_{13} h_{21} h_{22} h_{23} h_{31} h_{32} h_{33}

h_1 h_2 h_3

F norm says $h^T h = 1$

$$E(h) = \sum \|X_i \vec{h}\|^2 = \sum \vec{h}^T X_i^T X_i \vec{h}$$

$$= \vec{h}^T \left(\sum X_i^T X_i \right) \vec{h}$$

$\min E(\vec{h})$ s.t. $\vec{h}^T \vec{h} = 1$

you've seen this before!!

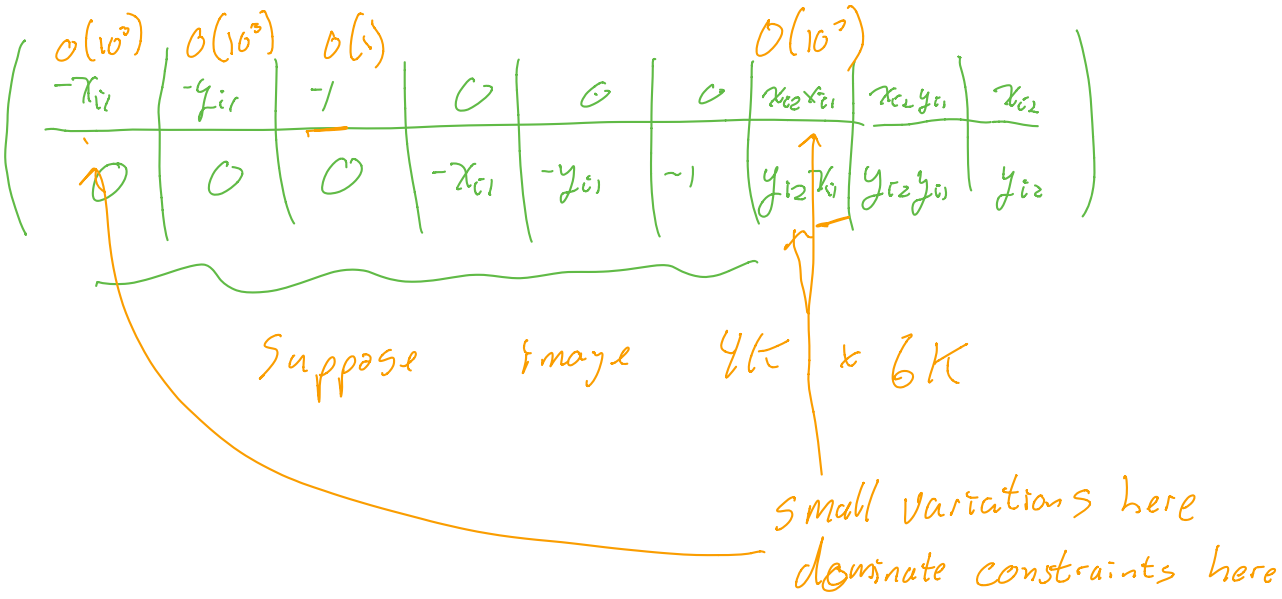
$\vec{h}$ is unit eigenvector corresponding to eigenvalue of $\sum X_i^T X_i$

solve using $SVD(\quad) = USV^T$

last row of V

Results are Disastrous

- Heavily skewed results, usually almost meaningless!
- Problem:
  - the terms of $\mathbf{h}_3$ are affected by the product of image coordinate values, whereas
  - the terms of $\mathbf{h}_1$ and $\mathbf{h}_2$ are only affected by the image coordinates individually, and so
  - errors in the image coordinates differentially affect the $\mathbf{h}_3$, which tend to be small to begin with...
- Solution is to "normalize" the constraints.

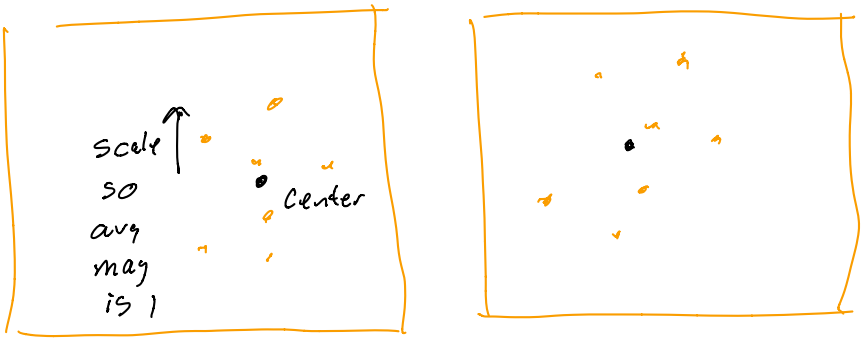


Normalization

Center and scale the pixel coordinates:

- In each image, compute the center of mass of the pixel coordinate vectors — the sets $\{\mathbf{x}_{i,1}\}$ and $\{\mathbf{x}_{i,2}\}$ — and shift the coordinate vectors by these centers.
- Scale each resulting set of coordinates so that the average magnitude of the pixel coordinate vectors (in each image separately) is 1.
- These centering and scaling can be described by affine transformation matrices, denoted $\mathbf{S}_1$ and $\mathbf{S}_2$, producing homogenous coordinates

$$\tilde{\mathbf{x}}'_{i,1} = \mathbf{S}_1\tilde{\mathbf{x}}_{i,1} \qquad \text{and} \qquad \tilde{\mathbf{x}}'_{i,2} = \mathbf{S}_2\tilde{\mathbf{x}}_{i,2}$$



$\mu_{x1} = \sum x_{i1}/N$

$\mu_{y1} = \sum y_{i1}/N$

$s_x = \sum |x_{i1} - \mu_{x1}|/N$

$s_y = \sum |y_{i1} - \mu_{y1}|/N$

same for image 2

S_1

$x'_{i1} = \dfrac{x_{i1} - \mu_{x1}}{s_x}$

$y'_{i1} = \dfrac{y_{i1} - \mu_{y1}}{s_y}$

$$\begin{pmatrix} x_{i1}' \\ y_{i1}' \\ 1 \end{pmatrix} = \begin{pmatrix} 1/s_{x1} & 0 & -\mu_{x1}/s_{x1} \\ 0 & 1/s_{y1} & -\mu_{y1}/s_{y1} \\ 0 & 0 & 1 \end{pmatrix} \begin{pmatrix} x_{i1} \\ y_{i1} \\ 1 \end{pmatrix}$$

$\tilde{x}'_{i1} = S_1 \tilde{x}_{i1}$

$\tilde{x}'_{i2} = S_2 \tilde{x}_{i2}$

Estimating the Homography

- Use the previous technique "algebraic distance" technique to solve for $\mathbf{H}'$.
- Gives the homography between the centered and normalized pixel values.
- Previous issue with the different types of error is mostly gone because of the normalization — errors are no longer magnified in different ways.
- We convert back to our unnormalized final estimate as

$$\mathbf{H} = \mathbf{S}_2^{-1}\mathbf{H}'\mathbf{S}_1$$

estimate H' using $\tilde{x}_{i1}'$ and $\tilde{x}_{i2}'$ values

$$\tilde{x}_{i2}' = H' \tilde{x}_{i1}'$$

$$S_2 \tilde{x}_{i2} = H' S_1 \tilde{x}_{i1}$$

$$\tilde{x}_{i2}' = \underbrace{S_2^{-1} H' S_1}_{H} \tilde{x}_{i1}$$

much much better

Discussion

Beware of bias!

- The foregoing solution works because we have eliminated most of the bias.
- Informally, bias is the disproportionate influence of a measured variable on the optimization, usually due to a non-linearity in the objective function.
- Nearly all computer vision problems have this bias.
- Normalization is one simple tool that can often be used to address bias. Sometimes more sophisticated tools are needed.

Solving the Geometric Distance Optimization

- Recall that the objective function is

$$\underbrace{\sum_i d(\tilde{\mathbf{x}}_{i,2}, \mathbf{H}\tilde{\mathbf{x}}_{i,1})^2}$$

  where we are back to using the geometric distance.

- We can think of this as a function $f(\mathbf{h})$, where $\mathbf{h}$ is formed from the nine entries in $\mathbf{H}$.

- We now need an iterative minimization technique, with the added constraint that $\mathbf{h}^\top \mathbf{h} = 1$.

- Initialized by the normalized estimated.

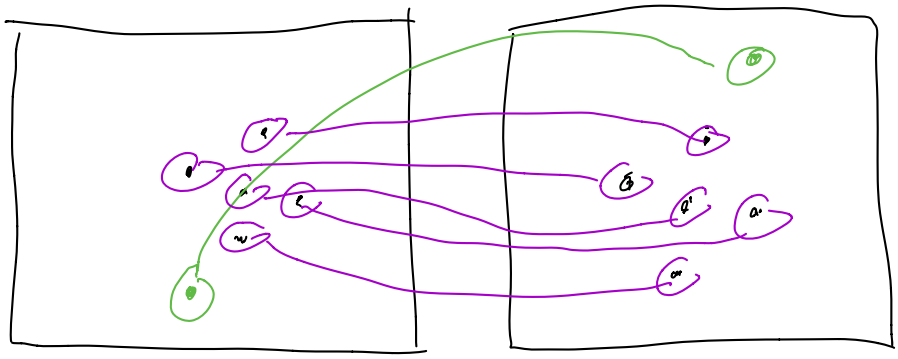
- Work in normalized coordinates(!)

- Solution is beyond our discussion.

Resetting the Stage

Subtitle: What about mismatches?

- One incorrect match can have a disastrous effect — worse even than for line estimation.
- Even with the ratio threshold of 0.8 from SIFT keypoint descriptor matches or the symmetric matching requirement used with ORB, we can not be assured of correct matches, especially for the harder image cases.
- Need a method for locating correct matches rather than throwing out bad matches after the estimation process has started.



Random Sampling — Outline

This is a restatement of the RANSAC algorithm in the setting of finding the best affine or homography transform. Much of it should be review.

1. Randomly choose a "minimal subset" of matches — enough to generate an estimate:
   - Four for the homography transformation
   - Three for the affine transformation
2. Generate an estimate from the minimal subset. Call it $\hat{\mathbf{H}}$.
3. Compute the error distances $d(\tilde{\mathbf{x}}_{i,2}, \hat{\mathbf{H}}\tilde{\mathbf{x}}_{i,1})$ for the remaining $N-4$ (or $N-3$) matches.
4. Evaluate the distances by either:
   (a) Counting the number whose error distance is less than a threshold (the "Ransac" method)
   (b) Computing an order statistic (such as the median) on the values.
5. **Repeat the foregoing** for some number, K, of randomly chosen minimal subsets, keeping the estimate and set of "inliers" that produces the best evaluation.
6. Apply the normalized, algebraic distance estimate to the inlier set and, optionally, compute a refined, non-linear estimate.

Generating the Minimal Subset

Somewhat different answers for the affine case and for the homography

- Affine: the least-squares solution becomes an exact solution (no error), when just 3 correspondences are involved.
- Homography:
  - Each correspondence yields two algebraic error terms:

$$\begin{aligned} x_{i,2}\tilde{\mathbf{x}}_{i,1}^\top \mathbf{h}_3 - \tilde{\mathbf{x}}_{i,1}^\top \mathbf{h}_1 = 0 \\ y_{i,2}\tilde{\mathbf{x}}_{i,1}^\top \mathbf{h}_3 - \tilde{\mathbf{x}}_{i,1}^\top \mathbf{h}_2 = 0 \end{aligned}$$

  - This may be written as $\mathbf{X}_i\mathbf{h} = \mathbf{0}$ where $\mathbf{X}_i$ is 2×9.
  - Stacking up four of these gives 8 constraints on $\mathbf{h}$.

$$\mathbf{X}\mathbf{h} = \mathbf{0}.$$

  - The ninth comes from the constraint that $\mathbf{h}^\top\mathbf{h} = 1$.
  - We find $\mathbf{h}$ as the unit basis vector for the right null space of $\mathbf{X}$. (We'll talk about how to do this in class.)

$X_i \vec{h}$ from earlier

2×9 X_i

$X_i \vec{h} = \vec{0}$

2 constraints on $\vec{h}$

$X = \begin{pmatrix} X_1 \\ X_2 \\ X_3 \\ X_4 \end{pmatrix}$

stack up constraints

$X\vec{h} = \vec{0}$

$\vec{h}^T\vec{h} = 1$

8×9

$\text{svd}(X) = U S V^T$

8×8 8×9 9×9

last row of V!!

$S = \begin{pmatrix} s_1 & & & 0 & 0 \\ & \ddots & & \vdots & \vdots \\ 0 & & s_8 & 0 & 0 \end{pmatrix}$

How Many Subsets are Needed?

Choosing the value of K from above

- Let p be the probability that one match is "good".
- Then p^4 is the probability that all four are good.
- $1 - p^4$ is the probability that at least one match is "bad"
- $(1 - p^4)^K$ is the probability that at least one match is bad in all K minimal subsets.
- $1 - (1 - p^4)^K$ is the probability that at least one minimal subset is good.
- Given p (determined empirically), we choose K to ensure that this probability reaches some threshold value (such as 0.99).

Degeneracies

- Affine: given a minimal set of three keypoint correspondences, all three keypoints from I_1 or all three from I_2 are colinear.
- Homography: given a minimal set of four correspondences, at least three keypoints from I_1 or at least three from I_2 are colinear.
- In practice we usually do not explicitly look for these cases because the computed transformation will be poor, having a large number of outliers, and therefore will be eliminated without needing a special test.

Summary: Solving the Two-Image Estimation Problem

1. Extract keypoints and descriptors from images I_1 and I_2.
2. Match keypoints based on matching descriptors and applying either the SIFT distance ratio test or the symmetric matching test.
3. Apply random sampling algorithm to select the "good matches"
4. Compute least-squares estimate based on the final set of good matches:
   - Use the geometric distance measure for the affine transformation.
   - Use the algebraic distance measure with normalized coordinates for the homography.

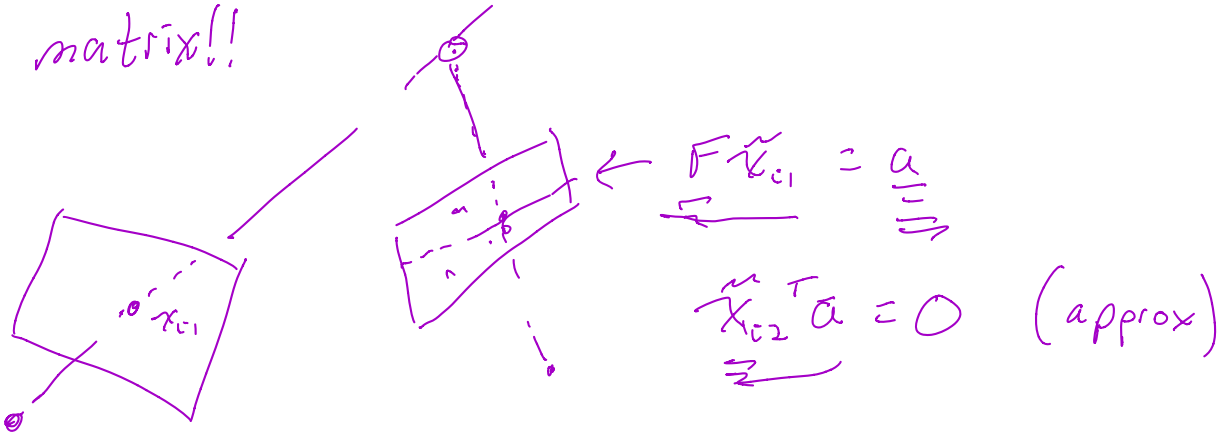
OpenCV provides tools. Use them!

Do the two image show same scene?

If yes Can the images be aligned (using $\tilde{H}$) well enough to form a clean mosaic (no ghosting)

If the images are same scene then there is an F matrix!!

Same scene



Good align if inliers from F most produce an accurate H

Multiple Image Estimation

We will briefly consider the following two questions:

- Which images overlap and therefore should have an **H** matrix computed for them?
- How do we align more than two images?

Which images?

It depends on keypoint matching

- Option 1:
  - Consider each pair I_i and I_j and apply keypoint matching
  - If there is a *sufficient* number of inlier matches to the final estimate, then consider I_i and I_j matched.

- Option 2:
  - Gather all keypoints into a spatial data structure and then match each keypoint from each image against the data structure.
  - When two images have more than a few matches between them, run the complete matching and estimation procedure.

Multiple Image Alignment

The correspondences are the key:

- Save the "good" keypoint matches for the matched image pairs
- Choose one image, call it I_0, to be the "anchor" on which to map the other images.
- Use correspondences to compute the mapping of all images onto I_0 simultaneously. If there are N images this produces $N-1$ homographies.

Unfortunately, we do not have time for the details.

After Estimation: Mapping the Images

We focus again on the two image case

- Suppose we have estimated the transformation from image I_1 to I_2 and now want to map image I_1 onto image I_2.
- Actually need to "inverse map", using the inverse transformation $\mathbf{H}^{-1}$.
- Start by forward mapping the image corners and computing an axis-aligned bounding box R on these corners.
- The union of this bounding box and the original I_2 is the combined image coordinate system.
- For each pixel location in R, inverse map using $\mathbf{H}^{-1}$ to find the pixel in I_1.
- If the pixel is inside I_1, then apply bilinear interpolation to compute its intensity (color) values.

Final Step: Blending

- Simplest technique is averaging.
- Can weight the intensities according to the distance from the center in order to obtain a smoother blend.
- There are also pyramid-based techniques.
- We can also estimate seams to avoid motion and parallax artifacts.

Summary

We have now taken an end-to-end tour of the mosaic application:

- Topics:
  - Image analysis
  - Keypoint and feature extraction
  - Camera modeling and transformations
  - Estimation
  - Image mapping
- Many of the issues and techniques we have discussed apply to a substantial number of other problems in computer vision.