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MIT CSAIL

6.869: Advances in Computer Vision



Lecture 13 Homographies and RANSAC

Depth-based ambiguity of position

Camera A Camera B



Under what conditions can you know where to translate each point of image A to where it would appear in camera B (with calibrated cameras), knowing nothing about image depths?



(a) camera rotation



and (b) imaging a planar surface



Geometry of perspective projection



Let's look at this scene from above...





Can generate any synthetic camera view as long as it has the same center of projection!









- When we only rotate the camera (around nodal point) depth does not matter
- It only performs a 2D warp
 - one-to-one mapping of the 2D plane
 - plus of course reveals stuff that was outside the field
 of view







• Now we just need to figure out this mapping

Aligning images: translation?





left on top





right on top



Translations are not enough to align the images



Homography



- Projective mapping between any two projection planes with the same center of projection
- called Homography
- represented as 3x3 matrix in homogenous coordinates

$$\begin{bmatrix} wx' \\ wy' \\ w \end{bmatrix} = \begin{bmatrix} * & * & * \\ * & * & * \\ * & * & * \end{bmatrix} \begin{bmatrix} x \\ y \\ I \end{bmatrix}$$

To apply a homography H

- Compute p' = Hp (regular matrix multiply)
- Convert p' from homogeneous to image coordinates (divide by w)



homography

Scamera rotates to here, same as Protating to here. $\pi_0 = A_0 \frac{P}{P}$ $\pi_1 = A_1 \frac{P}{P}$ 0,00 A. = - A.R A. = A. rotation matrix 0 0 0

homography

we seek Mis such that No = Mio Z, for all Xo X, A.P = Mio AoRP m Xo Xi for all P so X mult by R = (rotation) 0 Ao = Mio AoR ART = Mio Ao 1-000 AOR'B = Mio

homography



How many pairs of points does it take to specify M_10?

Planar objects





Figure 2.12 A point is projected into two images: (a) relationship between the 3D point coordinate (X, Y, Z, 1) and the 2D projected point (x, y, 1, d); (b) planar homography induced by points all lying on a common plane $\hat{n}_0 \cdot p + c_0 = 0$.

Mapping from one camera to another

What happens when we take two images of a 3D scene from different camera positions or orientations (Figure 2.12a)? Using the full rank 4×4 camera matrix $\tilde{P} = \tilde{K}E$ from (2.64), we can write the projection from world to screen coordinates as

$$\tilde{\boldsymbol{x}}_0 \sim \tilde{\boldsymbol{K}}_0 \boldsymbol{E}_0 \boldsymbol{p} = \tilde{\boldsymbol{P}}_0 \boldsymbol{p}. \tag{2.68}$$

Assuming that we know the z-buffer or disparity value d_0 for a pixel in one image, we can compute the 3D point location p using

$$p \sim E_0^{-1} \tilde{K}_0^{-1} \tilde{x}_0$$
 (2.69)

and then project it into another image yielding

$$\tilde{x}_1 \sim \tilde{K}_1 E_1 p = \tilde{K}_1 E_1 E_0^{-1} \tilde{K}_0^{-1} \tilde{x}_0 = \tilde{P}_1 \tilde{P}_0^{-1} \tilde{x}_0 = M_{10} \tilde{x}_0.$$
 (2.70)

Unfortunately, we do not usually have access to the depth coordinates of pixels in a regular photographic image. However, for a *planar scene*, as discussed above in (2.66), we can replace the last row of P_0 in (2.64) with a general *plane equation*, $\hat{n}_0 \cdot p + c_0$ that maps points on the plane to $d_0 = 0$ values (Figure 2.12b). Thus, if we set $d_0 = 0$, we can ignore the last column of M_{10} in (2.70) and also its last row, since we do not care about the final z-buffer depth. The mapping equation (2.70) thus reduces to

$$\tilde{x}_1 \sim \tilde{H}_{10} \tilde{x}_0$$
, (2.71)

where \tilde{H}_{10} is a general 3 × 3 homography matrix and \tilde{x}_1 and \tilde{x}_0 are now 2D homogeneous coordinates (i.e., 3-vectors) (Szeliski 1996). This justifies the use of the 8-parameter homography as a general alignment model for mosaics of planar scenes (Mann and Picard 1994; Szeliski 1996).

Images of planar objects, taken by generically offset cameras, are also related by a homography.

camera A



Measurements on planes



How to unwarp? CSE 576, Spring 2008

Projective Geometry

Image rectification



To unwarp (rectify) an image

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

CSE 576, Spring 2008

Solving for homographies

$$\begin{bmatrix} \mathbf{w} x_i' \\ \mathbf{w} y_i' \\ \mathbf{w} \end{bmatrix} \cong \begin{bmatrix} h_{00} & h_{01} & h_{02} \\ h_{10} & h_{11} & h_{12} \\ h_{20} & h_{21} & h_{22} \end{bmatrix} \begin{bmatrix} x_i \\ y_i \\ 1 \end{bmatrix}$$

$$x'_{i} = \frac{h_{00}x_{i} + h_{01}y_{i} + h_{02}}{h_{20}x_{i} + h_{21}y_{i} + h_{22}}$$
$$y'_{i} = \frac{h_{10}x_{i} + h_{11}y_{i} + h_{12}}{h_{20}x_{i} + h_{21}y_{i} + h_{22}}$$

$$\begin{aligned} x_i'(h_{20}x_i + h_{21}y_i + h_{22}) &= h_{00}x_i + h_{01}y_i + h_{02} \\ y_i'(h_{20}x_i + h_{21}y_i + h_{22}) &= h_{10}x_i + h_{11}y_i + h_{12} \end{aligned}$$

$$\begin{bmatrix} x_{i} & y_{i} & 1 & 0 & 0 & 0 & -x'_{i}x_{i} & -x'_{i}y_{i} & -x'_{i} \\ 0 & 0 & 0 & x_{i} & y_{i} & 1 & -y'_{i}x_{i} & -y'_{i}y_{i} & -y'_{i} \end{bmatrix} \begin{bmatrix} h_{00} \\ h_{01} \\ h_{02} \\ h_{10} \\ h_{11} \\ h_{12} \\ h_{20} \\ h_{21} \\ h_{22} \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \end{bmatrix}$$

CSE 576, Spring 2

Solving for homographies

Defines a least squares problem: minimize $||Ah - 0||^2$

- Since h is only defined up to scale, solve for unit vector ĥ
- Solution: \hat{h} = eigenvector of A^TA with smallest eigenvalue
- Works with 4 or more points

Image warping with homographies





automatic image mosaicing

- Basic Procedure
 - Take a sequence of images from the same position.
 - Rotate the camera about its optical center (entrance pupil).
 - Robustly compute the homography transformation between second image and first.
 - Transform (warp) the second image to overlap with first.
 - Blend the two together to create a mosaic.
 - If there are more images, repeat.

Robust feature matching through RANSAC



© Krister Parmstrand

Nikon D70. Stitched Panorama. The sky has been retouched. No other image manipulation.

with a lot of slides stolen from Steve Seitz and Rick Szeliski

15-463: Computational Photography Alexei Efros, CMU, Fall 2005

Feature matching



descriptors for left image feature points





descriptors for right image feature points



Strategies to match images robustly

(a) Working with individual features: For each feature point, find most similar point in other image (SIFT distance) Reject ambiguous matches where there are too many similar points

(b) <u>Working with all the features</u>: Given some good feature matches, look for possible homographies relating the two images

Reject homographies that don't have many feature matches.

(a) Feature-space outlier rejection

- Let's not match all features, but only these that have "similar enough" matches?
- How can we do it?
 - SSD(patch1,patch2) < threshold</p>
 - How to set threshold?
 Not so easy.



Feature-space outlier rejection

- A better way [Lowe, 1999]:
 - 1-NN: SSD of the closest match
 - 2-NN: SSD of the second-closest match
 - Look at how much better 1-NN is than 2-NN, e.g. 1-NN/2-NN
 - That is, is our best match so much better than the rest?



Feature-space outlier rejection



- Can we now compute H from the blue points?
 - No! Still too many outliers...
 - What can we do?

(b) Matching many features--looking for a good homography

Simplified illustration with translation instead of homography



Note: at this point we don't know which ones are good/bad





0 inliers



4 inliers



Select one match, count inliers

Keep match with largest set of inliers

At the end: Least squares fit



Find "average" translation vector, but with only inliers

Reference

- M. A. Fischler, R. C. Bolles. Random Sample Consensus: A Paradigm for Model Fitting with Applications to Image Analysis and Automated Cartography. Comm. of the ACM, Vol 24, pp 381-395, 1981.
- <u>http://portal.acm.org/</u> <u>citation.cfm?id=358692</u>

Graphies and Inage Processing I. D. Foley Editor Random Sample Consensus: A Paradigm for Model Fitting with Applications to Image Analysis and Automated Cartography

Martin A. Fischler and Robert C. Bolles SRI International

A new paradigm, Random Sample Consensus (RANSAC), for fitting a model to experimental data is introduced. RANSAC is capable of interpreting/ smoothing data containing a significant percentage of gross errors, and is thus ideally suited for applications in automated image analysis where interpretation is based on the data provided by error-prone feature detectors. A major portion of this paper describes the application of RANSAC to the Location Determination Problem (LDP): Given an image depicting a set of landmarks with known locations, determine that point in space from which the image was obtained. In response to a RANSAC requirement, new results are derived on the minimum number of landmarks needed to obtain a solution, and algorithms are presented for computing these minimum-landmark solutions in closed form. These results provide the basis for an automatic system that can solve the LDP under difficult viewing

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and analysis conditions. Implementation details and computational examples are also presented. Key Words and Phrases: model fitting, scene analysis, camera calibration, image matching, location determination, automatof cartography. CR Categories: 360, 361, 371, 50, 81, 8.2

I. Introduction

We introduce a new paradigm, Random Sample Consensus (RANSAC), for fitting a model to experimental data; and illustrate its use in scene analysis and automated cartography. The application discussed, the location determination problem (LDP), is treated at a level beyond that of a mere example of the use of the RANSAC paradigm; new basic findings concerning the conditions under which the LDP can be solved are presented and a comprehensive approach to the solution of this problem that we anticipate will have near-term practical applications is described.

To a large extent, scene analysis (and, in fact, science in general) is concerned with the interpretation of sensed data in terms of a set of predefined models. Conceptually, interpretation involves two distinct activities: First, there is the problem of finding the best match between the data and one of the available models (the classification problem); Second, there is the problem of computing the best values for the free parameters of the selected model (the parameter estimation problem). In practice, these two problems are not independent—a solution to the parameter estimation problem is often required to solve the classification problem.

Classical techniques for parameter estimation, such as least squares, optimize (according to a specified objective function) the fit of a functional description (model) to all of the presented data. These techniques have no internal mechanisms for detecting and rejecting gross errors. They are averaging techniques that rely on the assumption (the smoothing assumption) that the maximum expected deviation of any datum from the assumed model is a direct function of the size of the data set, and thus regardless of the size of the data set, there will always be enough good values to smooth out any gross deviations.

In many practical parameter estimation problems the smoothing assumption does not hold; i.e., the data contain uncompensated gross errors. To deal with this situation, several heuristics have been proposed. The technique usually employed is some variation of first using all the data to derive the model parameters, then locating the datum that is farthest from agreement with the instantiated model, assuming that it is a gross error, deleting it, and iterating this process until either the maximum deviation is less then some preset threshold or until there is no longer sufficient data to proceed.

It can easily be shown that a single gross error ("poisoned point"), mixed in with a set of good data, can

Communications of the ACM	June 1981 Volume 24 Number 6

RANSAC for estimating homography

- RANSAC loop:
- Select four feature pairs (at random)
- Compute homography H (exact)
- Compute inliers where $||p_i'|$, $H p_i || < \epsilon$
- Keep largest set of inliers
- Re-compute least-squares H estimate using all of the inliers

Simple example: fit a line

• Rather than homography H (8 numbers) fit y=ax+b (2 numbers a, b) to 2D pairs


- Pick 2 points
- Fit line
- Count inliers

- Pick 2 points
- Fit line



- Pick 2 points
- Fit line
- Count inliers 9 inlier • • • • •

- Pick 2 points
- Fit line
- Count inliers 8 inlier • • • • •

- Use biggest set of inliers
- Do least-square fit



RANSAC





red:

rejected by 2nd nearest neighbor criterion blue: Ransac outliers yellow:





Robustness

- Proportion of inliers in our pairs is G (for "good")
- Our model needs P pairs
 - P=4 for homography
- Probability that we pick P inliers? $-G^{P}$
- Probability that after N RANSAC iterations we have not picked a set of inliers? $-(1-G^{P})^{N}$

Robustness: example

- Matlab: p=4; x=0.5; n=1000; (1-x^p)^n
- Proportion of inliers G=0.5
- Probability that we pick P=4 inliers?
 0.5⁴=0.0625 (6% chance)
- Probability that we have not picked a set of inliers?
 - N=100 iterations:

 $(1-0.5^4)^{100}=0.00157$ (1 chance in 600)

- -N=1000 iterations:
 - 1 chance in 1e28

Robustness: example

• Proportion of inliers G=0.3



- Probability that we pick P=4 inliers?
 0.3⁴=0.0081 (0.8% chance)
- Probability that we have not picked a set of inliers?
 - N=100 iterations:

 $(1-0.3^4)^{100}=0.44$ (1 chance in 2)

- -N=1000 iterations:
 - 1 chance in 3400

Robustness: example

• Proportion of inliers G=0.1



- Probability that we pick P=4 inliers?
 0.1⁴=0.0001 (0.01% chances, 1 in 10,000)
- Probability that we have not picked a set of inliers?
 - N=100 iterations: (1-0.1⁴)¹⁰⁰=0.99
 - N=1000 iterations: 90%
 - N=10,000: 36%
 - N=100,000: 1 in 22,000

Robustness: conclusions

- Effect of number of parameters of model/ number of necessary pairs
 - Bad exponential
- Effect of percentage of inliers
 - Base of the exponential
- Effect of number of iterations
 - Good exponential

RANSAC recap

- For fitting a model with low number P of parameters (8 for homographies)
- Loop
 - Select P random data points
 - Fit model
 - Count inliers

(other data points well fit by this model)

• Keep model with largest number of inliers

Example: Recognising Panoramas

M. Brown and D. Lowe, University of British Columbia

* M. Brown and D. Lowe. Automatic Panoramic Image Stitching using Invariant Features. International Journal of Computer Vision, 74(1), pages 59-73, 2007 (pdf 3.5Mb | bib) * M. Brown and D. G. Lowe. Recognising Panoramas. In Proceedings of the 9th International Conference on Computer Vision (ICCV2003), pages 1218-1225, Nice, France, 2003 (pdf 820kb | ppt | bib)

"Recognising Panoramas"?











RANSAC for Homography



RANSAC for Homography



RANSAC for Homography





























Results





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Serratus

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Benefits of Laplacian image compositing



(a) Linear blending



(b) Multi-band blending

Figure 7. Comparison of linear and multi-band blending. The image on the right was blended using multi-band blending using 5 bands and $\sigma = 5$ pixels. The image on the left was linearly blended. In this case matches on the moving person have caused small misregistrations between the images, which cause blurring in the linearly blended result, but the multi-band blended image is clear.

M. Brown and D. Lowe. Automatic Panoramic Image Stitching using Invariant Features. International Journal of Computer Vision, 74(1), pages 59-73, 2007

Photo Tourism: Exploring Photo Collections in 3D

Noah Snavely Steven M. Seitz University of Washington Richard Szeliski Microsoft Research

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Photo Tourism Exploring photo collections in 3D

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SIGGRAPH 2006

Photo Tourism overview



Photo Tourism overview



Scene reconstruction

- Automatically estimate
 - position, orientation, and focal length of cameras
 - 3D positions of feature points



Feature detection

Detect features using SIFT [Lowe, IJCV 2004]





Detect features using SIFT [Lowe, IJCV 2004]



Feature detection

Detect features using SIFT [Lowe, IJCV 2004]



Feature matching

Match features between each pair of images



Feature matching

Refine matching using RANSAC [Fischler & Bolles 1987] to estimate fundamental matrices between pairs

(See 6.801/6.866 for fundamental matrix, or Hartley and Zisserman, Multi-View Geometry. See also the fundamental matrix song: http://danielwedge.com/fmatrix/)



Structure from motion



Links

- Code available: <u>http://phototour.cs.washington.edu/bundler/</u>
- http://phototour.cs.washington.edu/
- http://livelabs.com/photosynth/
- http://www.cs.cornell.edu/~snavely/