EE5490: Image Signal Processing Lab-4

Image mosaicing

Feb. 17 (Batch-A) and Feb. 21 (Batch-B)

1 Problem statement

Image mosaicing is the alignment and stitching of a collection of images having overlapping regions into a single image. In this assignment, you have been given three images which were captured by panning the scene left to right. These images (img1.pgm, img2.pgm and img3.pgm) capture overlapping regions of the same scene from different viewpoints. The task is to determine the geometric transformations (homographies) between these images and stitch them into a single image.

2 Steps

- 1. Take img2.pgm as the reference image.
- 2. Determine homography H_{21} between $I_2 = \text{img2.pgm}$ and $I_1 = \text{img1.pgm}$ such that $I_1 = H_{21}I_2$.
- 3. Determine homography H_{23} between $I_2=\text{img2.pgm}$ and $I_3=\text{img3.pgm}$ such that $I_3=H_{23}I_2$.
- 4. Create an empty canvas. For every pixel in the canvas, find corresponding points in I_1 , I_2 and I_3 using H_{21} , identity matrix and H_{23} respectively (target-to-source mapping). Blend the three values by averaging them. Employ the values in blending only if it falls within the corresponding image bounds. Choose the origin so as to get a full mosaic.

2.1 Determining homography between two images

1. Determine SIFT features of the two images and determine correspondences between them. File sift_corresp.m returns the SIFT correspondences between two images (see Section 2.3). Now to find H such that:

$$\begin{bmatrix} x_i' \\ y_i' \\ 1 \end{bmatrix} \sim H \begin{bmatrix} x_i \\ y_i \\ 1 \end{bmatrix}$$

- 2. Run RANSAC on matched points (correspondences) to remove outliers (wrong matches), and find the homography between the two images.
 - (a) Input: Matched points (x_i, y_i) and (x_i', y_i') with $i \in \mathcal{M}$.
 - (b) Randomly pick four correspondences (so that we can form eight equations), i.e. (x_i, y_i) and (x_i', y_i') with $i \in \mathcal{R} \subset \mathcal{M}$ and $|\mathcal{R}| = 4$, where $|\cdot|$ denotes the cardinality of the set.
 - (c) Calculate the homography H using the above four correspondences (see Section 2.2).
 - (d) For each of the remaining correspondences (x_i, y_i) and (x_i', y_i') with $i \in \mathcal{P} = \mathcal{M} \setminus \mathcal{R}$, check whether they satisfy the homography (within an error bound). If yes, add the index of that correspondence to the consensus set.

$$\begin{bmatrix} x_i'' \\ y_i'' \\ z_i'' \end{bmatrix} \leftarrow H \begin{bmatrix} x_i \\ y_i \\ 1 \end{bmatrix}, \text{ and normalize so that } z_i'' = 1,$$
 i.e. $x_i'' \leftarrow x_i''/z_i''$ and $y_i'' \leftarrow y_i''/z_i''$

If
$$\sqrt{(x_i'-x_i'')^2-(y_i'-y_i'')^2}<\varepsilon (=10)$$
, then update consensus set $\mathcal{C}\leftarrow\mathcal{C}\cup\{i\}$.

- (e) If the consensus set is large enough i.e. if $|\mathcal{C}| > d (= 0.8|\mathcal{P}|)$, then return this homography H, else go to step (b).
- (f) Output: Homography H.

2.2 Calculating homography

1. Consider a correspondence (x, y) and (x', y'),

$$\begin{bmatrix} x' \\ y' \\ z' \end{bmatrix} = \begin{bmatrix} h_1 & h_2 & h_3 \\ h_4 & h_5 & h_6 \\ h_7 & h_8 & h_9 \end{bmatrix} \begin{bmatrix} x \\ y \\ 1 \end{bmatrix}.$$

Upon normalizing z',

$$x' = h_1x + h_2y + h_3/h_7x + h_8y + h_9,$$

 $y' = h_4x + h_5y + h_6/h_7x + h_8y + h_9.$

Form two equations for each correspondence (corresponding to two rows of matrix A).

$$\begin{split} (x)h_1 + (y)h_2 + (1)h_3 + (0)h_4 + (0)h_5 + (0)h_6 \\ - (x'x)h_7 - (x'y)h_8 - (x')h_9 &= 0 \\ (0)h_1 + (0)h_2 + (0)h_3 + (x)h_4 + (y)h_5 + (1)h_6 \\ - (y'x)h_7 - (y'y)h_8 - (y')h_9 &= 0 \end{split}$$

2. Solve the system,

$$A_{8\times 9} \begin{bmatrix} h_1 \\ h_2 \\ \vdots \\ h_9 \end{bmatrix} = 0$$

i.e., find the null space of A.

3. Homography matrix

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

Normalize H so that $h_9=1$.

2.3 Using SIFT

Files needed:

- 1. sift_corresp.m
- 2. sift.m and
- 3. sift (for GNU/Linux) or siftWin32.exe (for Windows).

Copy the above three files to the working directory. In GNU/Linux, check that the file sift has executable permission. If not, run chmod +x sift in a terminal.

Usage: [corresp1, corresp2] = sift_corresp('img1.pgm','img2.pgm')

-end-