

ECE 661 Fall 2018

Homework #5

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1 Methodology

In the homework, I select the SURF algorithm to find and match the interest points. The LSM in Sec. 1.1 is applied in RANSAC, Sec. 1.2, to find inliners. Levenberg-Marquardt Algorithm, Sec. 1.3, is applied to find homography by the inliners. For Monsaicing image, please see Sec. 1.4.

1.1 Least Square Method (LSM) by Singular Value Decomposition (SVD)

Assuming $x' = Hx$ that x and x' are expressed using HC:

$$x = \begin{bmatrix} x_1 \\ x_2 \\ x_3 \end{bmatrix} \text{ and } x' = \begin{bmatrix} x'_1 \\ x'_2 \\ x'_3 \end{bmatrix}$$

with

$$H = \begin{bmatrix} h_{11} & h_{12} & h_{13} \\ h_{21} & h_{22} & h_{23} \\ h_{31} & h_{32} & h_{33} \end{bmatrix}.$$

We end up with

$$\begin{cases} x'_1 = h_{11}x_1 + h_{12}x_2 + h_{13}x_3 \\ x'_2 = h_{21}x_1 + h_{22}x_2 + h_{23}x_3 \\ x'_3 = h_{31}x_1 + h_{32}x_2 + h_{33}x_3 \end{cases}$$

Since x' is in HC, the 2D physical coordinates (x', y') have

$$\begin{cases} x' = \frac{x'_1}{x'_3} = \frac{h_{11}x_1 + h_{12}x_2 + h_{13}x_3}{h_{31}x_1 + h_{32}x_2 + h_{33}x_3} \\ y' = \frac{x'_2}{x'_3} = \frac{h_{21}x_1 + h_{22}x_2 + h_{23}x_3}{h_{31}x_1 + h_{32}x_2 + h_{33}x_3} \end{cases}.$$

Since there is no ideal points in real image, $x' \neq 0$, we have

$$\begin{cases} h_{11}x_1x'_3 + h_{12}x_2x'_3 + h_{13}x_3x'_3 - h_{31}x_1x'_1 - h_{32}x_2x'_1 - h_{33}x_3x'_1 = 0 \\ h_{21}x_1x'_3 + h_{22}x_2x'_3 + h_{23}x_3x'_3 - h_{31}x_1x'_2 - h_{32}x_2x'_2 - h_{33}x_3x'_2 = 0 \end{cases}.$$

While we select the points on real images, $x_3 = 1$ and $x'_3 = 1$. And H in HC, we can assign $h_{33} = 1$. We then have

$$\begin{cases} h_{11}x_1 + h_{12}x_2 + h_{13} - h_{31}x_1x'_1 - h_{32}x_2x'_1 - x'_1 = 0 \\ h_{21}x_1 + h_{22}x_2 + h_{23} - h_{31}x_1x'_2 - h_{32}x_2x'_2 - x'_2 = 0 \end{cases}.$$

We then formulate it into a matrix expression with a vector

$$\vec{H} = [h_{11}, h_{12}, h_{13}, h_{21}, h_{22}, h_{23}, h_{31}, h_{32}, h_{33}]^T$$

$$A\vec{H} = 0, \quad (1)$$

where

$${}^{2i \times 9} A = \begin{bmatrix} x_1^{(1)} & x_2^{(1)} & 1 & 0 & 0 & 0 & -x_1^{(1)}x_1'^{(1)} & -x_2^{(1)}x_1'^{(1)} & -x_1'^{(1)} \\ 0 & 0 & 0 & x_1^{(1)} & x_2^{(1)} & 1 & -x_1^{(1)}x_2'^{(1)} & -x_2^{(1)} & -x_2'^{(1)} \\ \dots & \dots \\ \dots & \dots \\ x_1^{(i)} & x_2^{(i)} & 1 & 0 & 0 & 0 & -x_1^{(i)}x_1'^{(i)} & -x_2^{(i)}x_1'^{(i)} & -x_1'^{(i)} \\ 0 & 0 & 0 & x_1^{(i)} & x_2^{(i)} & 1 & -x_1^{(i)}x_2'^{(i)} & -x_2^{(i)}x_2'^{(i)} & -x_2'^{(i)} \end{bmatrix}$$

We can solve this by SVD, which gives $A = UDV^H$. Please note, the number i of corresponding pairs needs to be greater than 5 in order to satisfy the rank of A is greater than 9, otherwise we cannot apply SVD on A . In my application, I set $i = 9$. The last column of V is \vec{H} , which is also corresponding to the minimum value of D , which is actually the minimal variance. We then reshape \vec{H} to the matrix H .

1.2 Random Sample Consensus (RANSAC)

RANSAC is an algorithm based on probability to remove the bad samples. In our homework, it removes the outliers matched by SURF features. There are some parameters listed as follows,

- δ - decision threshold. If the distance d between the estimated point \tilde{x}' and the real target point x' greater than δ , we will regard it as a outlier. I set $\delta = 20$ in chess board distance.
- ϵ - the proportion of outliers to all pairs. I set $\epsilon = 0.2$.
- p - the probability that at least one of the N trials will be free of outliers. I set $p = 0.99$.
- N - number of trails,

$$N = \frac{\ln(1-p)}{\ln(1-(1-\epsilon)^m)}$$

- m - the set of correspondences for calculating homography matrix H in each trail. I set $m = 9$, which is actually the i in Sec. 1.1.
- M - a minimum value for the size of the inlier set for it to be acceptable,

$$M = (1-\epsilon)n_{total}$$

- n_{total} - total number of SURF matched pairs.

The RANSAC is explained as following steps.

1. Randomly sample m corresponding pairs, then use LSM to find the homography.
2. Transform each point x_i by the homography and get the estimated point \tilde{x}'_i . Compute the distance between and x'_i which is

$$d_i = \text{sum}(\text{abs}(\tilde{x}'_i - x'_i))$$

3. Count the number of $d_i < \delta$, if it is grater than M , then we mark these pairs as a candidate inlier set.
4. Repeat 1st step N times, and chose the set with largest number of inlier set as the inliers.

The final survive set are inliers, and we are going to use LM Algorithm to find the homography from this set in next Sec. 1.3.

1.3 Levenberg-Marquardt Algorithm (LM)

LM algorithm is nonlinear optimization method based on least squares. Assume the matched points x_i in the image A forms a set as X , the corrsponding matched points x'_i in the image B is X' , where $X' = HX$. Please note, here we use Homogeneous Coordinates. Then we want to find a minimizer H , where

$$\tilde{H} = \underset{H}{\text{argmin}} (X' - HX)^2,$$

This is the problem we want to solve. The we regard H as a vector \vec{H} as we stated in Sec. 1.1. We formulated a function $f(\vec{H}) = HX$ for notation convinence. Then the problem is re-stated as

$$\tilde{H} = \underset{H}{\text{argmin}} (X' - f(\vec{H}))^2$$

To start a minimization, the user has to provide an initial guess for the parameter vector \vec{H}_0 . Usually, I like to start with $H_0 = [1, \dots, 1]$.

The Jacobina of $f(\vec{H})$ is

$${}^{3 \times 9} J_f = \begin{bmatrix} \frac{\partial f_1}{\partial h_{11}} & \dots & \frac{\partial f_1}{\partial h_{33}} \\ \frac{\partial f_2}{\partial h_{11}} & \dots & \frac{\partial f_2}{\partial h_{33}} \\ \frac{\partial f_3}{\partial h_{11}} & \dots & \frac{\partial f_3}{\partial h_{33}} \end{bmatrix}.$$

Please recall 1st order Taylor expansion,

$$f(\vec{H} + \vec{\delta}) \approx f(\vec{H}) + \epsilon(\vec{H}) = f(\vec{H}) + J_f \vec{\delta},$$

where $\epsilon(\vec{H}) = f(\vec{H} + \vec{\delta}) - f(\vec{H})$.

Similar as Gaussian-Newton method, we need to find the best $\vec{\delta}$ for getting directly to the minimum.

$$(J_f^T J_f) \vec{\delta} = J_f^T \epsilon(\vec{H})$$

Till here all above are the same as Gaussian-Newton method. Focusing on the fact that, if $J_f^T J_f$ is purely diagonal, we add a damper as μI , I is an identity matrix. Therefore we have

$$\begin{aligned} (J_f^T J_f + \mu I) \vec{\delta} &= J_f^T \epsilon(\vec{H}) \\ \Rightarrow \vec{\delta} &= (J_f^T J_f + \mu I)^{-1} J_f^T \epsilon(\vec{H}) \\ \Rightarrow \tilde{\vec{H}} &= \tilde{\vec{H}_0} + \vec{\delta} \end{aligned}$$

In practice, it is impossible to get $\tilde{\vec{H}}$ in one step. You may run several iterations from above equations to find the optima.

1.4 Mosaicing Method

The Mosaicing is to stitch together a sequence of overlapping photos of a scene to create a single panoramic photo. In the homework, we are required to use pairwise homographies to project all images into the coordinate frame of the central image in the sequence.

Since we have done 2 images projection in the homework 2, the only problem is that after each 1-by-1 projection, the coordinate frame of the new image will change. The homography will not match to the new image. The problem can be easily solved by the following method.

We assume all images combining are the same size $h \times w$, and each sequential pair has the homography H_i , where $x_{i+1} = H_i x_i$. The center image is I_c .

1. Calculate all Homography from each image I_i to the central image I_c .
 - For left side images ($c > i$), $H_{i \rightarrow c} = \prod_{j=i}^{j < c} H_j$.
 - For right side images ($c < i$), $H_{i \rightarrow c} = (\prod_{j=c}^{j < i} H_j)^{-1}$.
2. Use the corner points in the images $(0,0)$, $(h,0)$, $(0,w)$ and (h,w) to project by each $H_{i \rightarrow c}$ and get new corner points $(h_i(k), w_i(k))$, where $k = 1, 2, 3, 4$ corresponding sequentially to the projection of $(0,0)$, $(h,0)$, $(0,w)$ and (h,w) .
3. Find the minimum and maximum separately of $h_i(k)$ and $w_i(k)$ as $\min(h_i(k))$, $\max(h_i(k))$, $\min(w_i(k))$ and $\max(w_i(k))$.
4. The Mosaicing image size is $(\max(h_i(k)) - \min(h_i(k))) \times (\max(w_i(k)) - \min(w_i(k)))$; There is an offset for every image is $\min(h_i(k))$ and $\min(w_i(k))$. You can change either manually add the offset while projecting, or change the homographies by the following steps,

- (a) Standardize the homography

$$H_{i \rightarrow \text{New}} = \frac{H_{i \rightarrow c}}{H_{i \rightarrow c}(3,3)}$$

- (b) Add the offset on height

$$H_{i \rightarrow \text{New}}(1,3) = H_{i \rightarrow c}(1,3) + \min(h_i(k))$$

- (c) Add the offset on width

$$H_{i \rightarrow \text{New}}(2,3) = H_{i \rightarrow c}(2,3) + \min(w_i(k))$$

- (d) Project all images into the new Mosaicing image frame by $x_{\text{New}} = H_{i \rightarrow \text{New}} x_i$

2 Test Picture Set

The following 5 pictures are the original pictures for this homework. The pictures are taken sequentially from left to right by a iPhone 7.



(a) 1.jpg



(b) 2.jpg



(c) 3.jpg



(d) 4.jpg



(e) 5.jpg

Figure 1: My picture set for homework.

3 Results

3.1 SURF Match and RANSAC

The blue cross, \times , represents the interest points on the left images. The red circle, \circ , represents the interest points on the right images. The green line represents the [inlier](#) matching pairs. The blue line represents the [outlier](#) matching pairs.



(a) SURF match between 1.jpg and 2.jpg



(b) SURF match between 2.jpg and 3.jpg



(c) SURF match between 3.jpg and 4.jpg

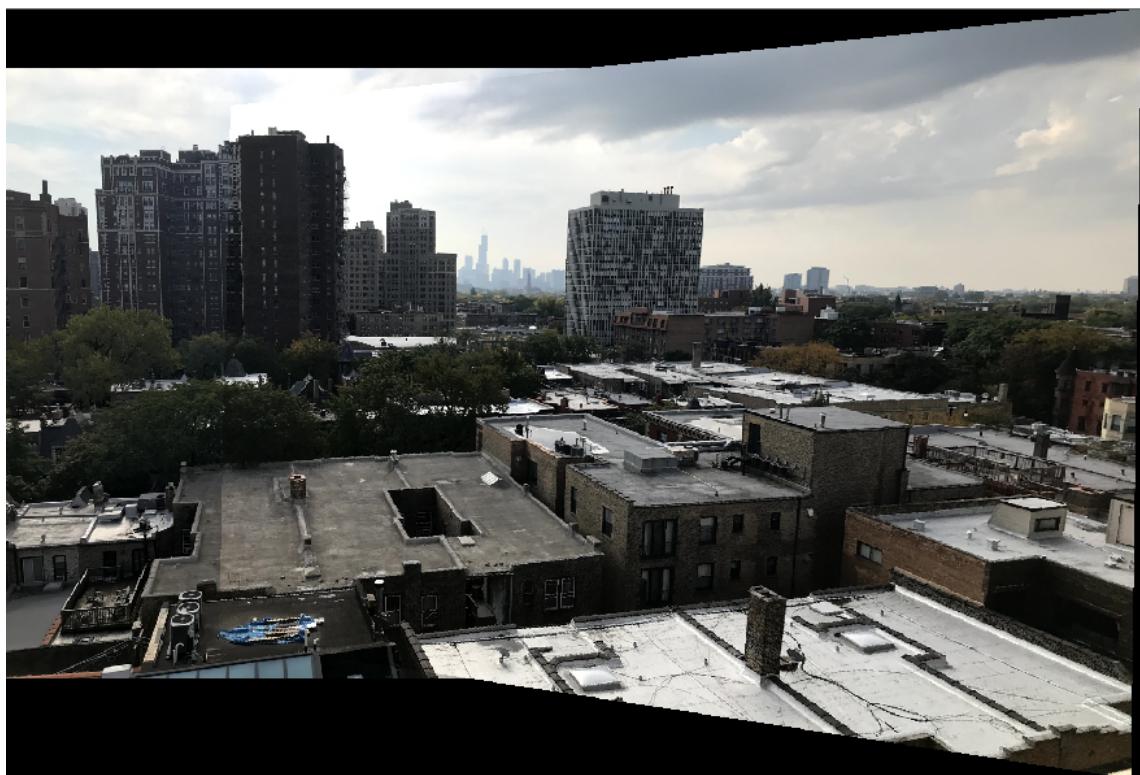


(d) SURF match between 4.jpg and 5.jpg

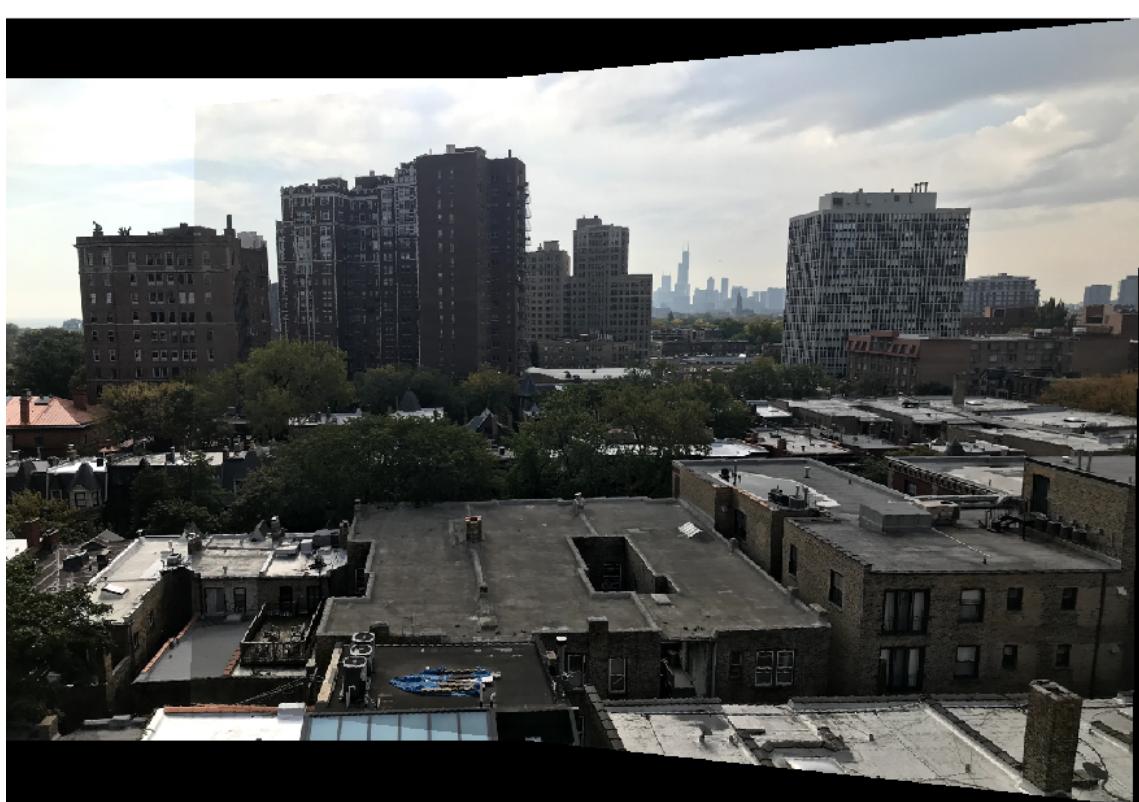
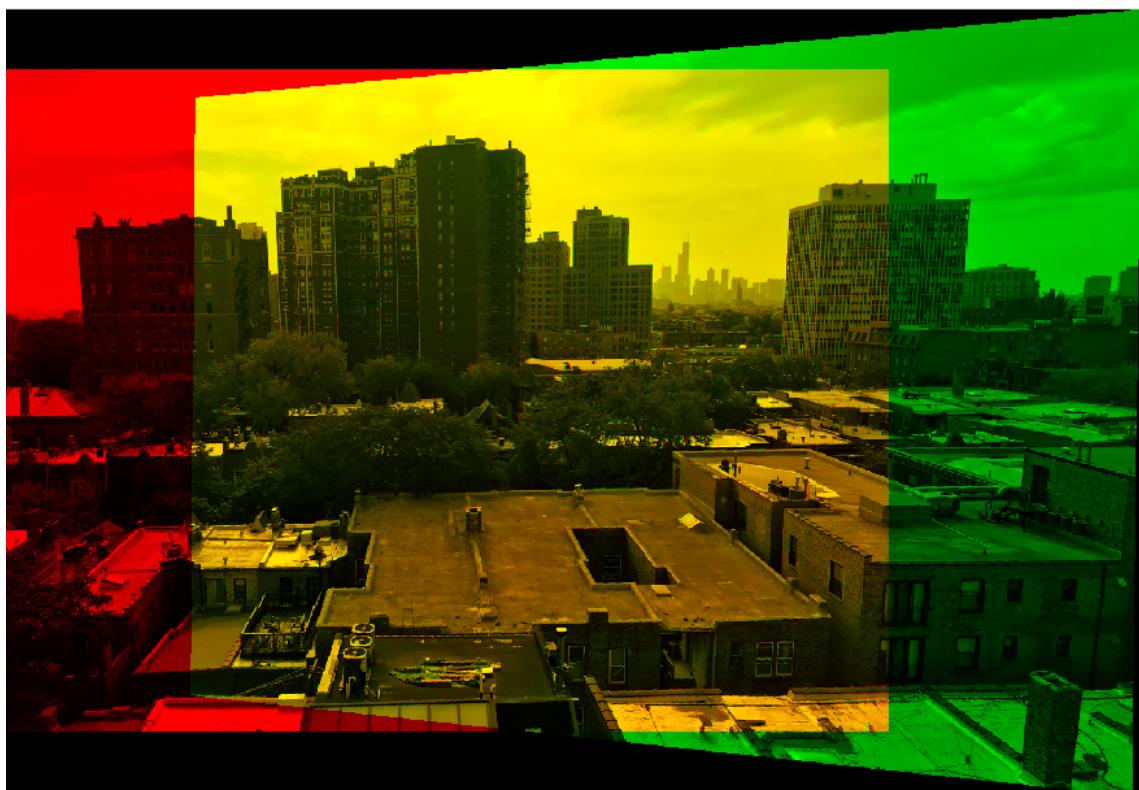
Figure 2: The SURF match results after RANSAC algorithm.

3.2 Projection by Homography from AuHoCa()

The following pictures are generate by the transformation according to the pairwise homographies calculated by the AuHoCa function. Please notice, for better visualization, the projecting picture is tinted in **green**; the projected picture is tinted in **red**; their intersection is tinted in **yellow** due to the overlapping. There is an un-tinted picture shown below each tinted picture.



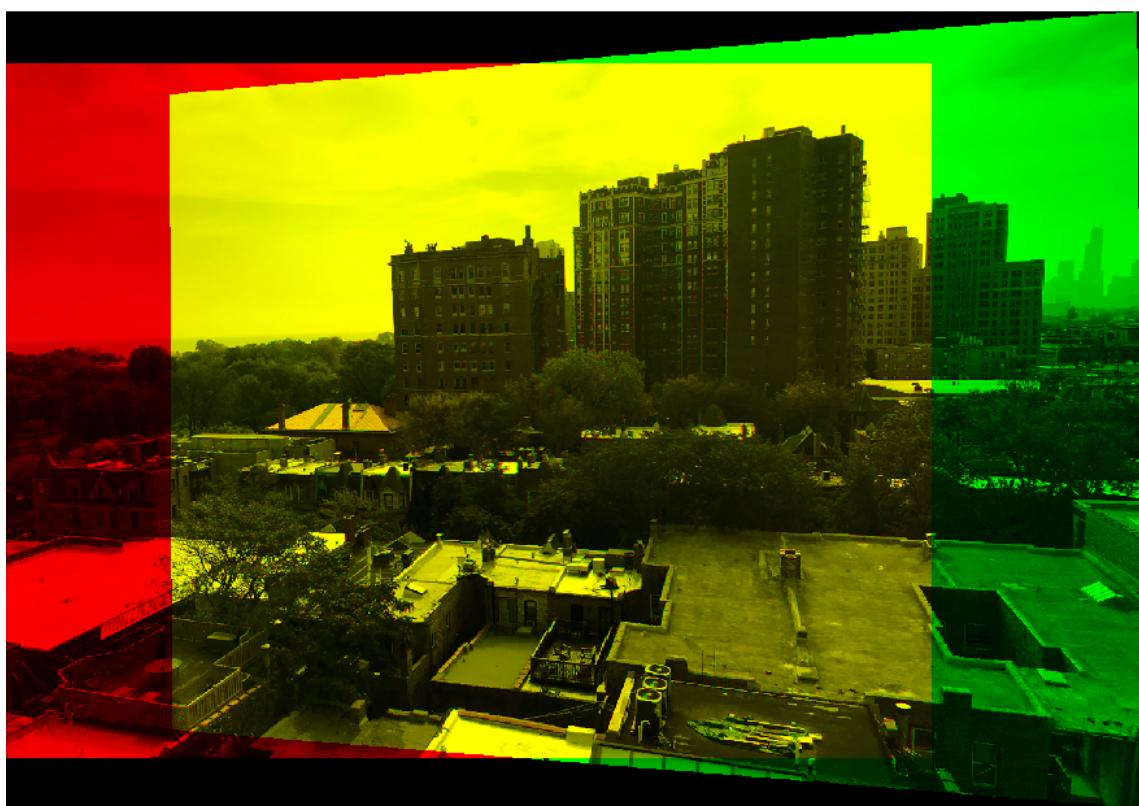
(a) Projection of 1.jpg on 2.jpg



(b) Projection of 2.jpg on 3.jpg



(c) Projection of 3.jpg on 4.jpg

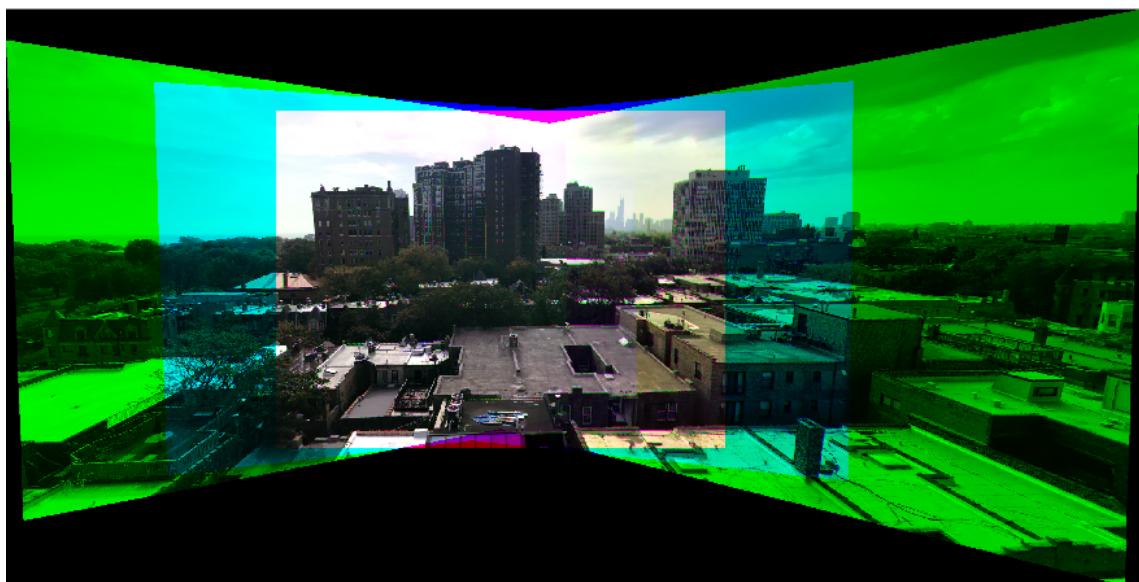


(d) Projection of 4.jpg on 5.jpg

Figure 3: The projection results by AuHoCa Homographies.

3.3 Monsaicing Image

The Monsaicing image Fig. 4 is generate by the transformation according to the pairwise homographies calculated by the AuHoCa function. In upper figure, Fig. 4a, the 3.jpg is tinted in red as the base image; the 1.jpg and 5.jpg is tinted in green; the 2.jpg and 4.jpg is tinted in blue. Due to the overlapping, the color varies at the intersection. There is an un-tinted picture Fig. 4b shown after tinted picture.



(a) The tinted Monsaicing image



(b) The un-tinted Monsaicing image

Figure 4: The final Monsaicing image.

4 Source Code

The code is written in Matlab 2018b and Windows 10.

4.1 Main Script

```

1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi349@purdue.edu
4
5 close all;clear; clc;
6
7 %% read all images
8 img = cell(5,1);
9 for i = 1:5
10     img{i} = imresize(imread([pwd , '\MyPics\ ', num2str(i) , '.jpg']), [900,1200]);
11 % img{i} = imread([pwd , '\MyPics\ ', num2str(i) , '.jpg']);
12 end
13
14 %% Auto Homography Calculation
15 H = cell(4,1);
16 Newimg = cell(4,1);
17 for i = 1:4 % i=1 corresponds to 1.jpg to 2.jpg
18     H{i} = AuHoCa(img{i},img{i+1},[num2str(i) , 'To' , num2str(i+1)]);
19     projectH(img{i},img{i+1},H{i},1,[num2str(i) , 'To' , num2str(i+1)]);
20     Newimg{i} = projectH(img{i},img{i+1},H{i},0,[num2str(i) , 'To' , num2str(i+1)]);
21 end
22
23 %% Mosaicing 5 pictures
24 Finalimg = mosaicing5(img,H,[ 'SuperFinal' ]);

```

4.2 Automatic Homography Calculator

```
1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi349@purdue.edu
4
5 function H = AuHoCa(img1,img2,filename)
6 % Automatic Homography Calculator
7 % Find matched SURF features pairs
8 if nargin <3
9     filename = datetime('today');
10 end
11
12 pairs = surfMatch(img1,img2,filename);
13 % RANSAC
14 [bestIdx,~] = ransacH(pairs);
15
16
17 pltPairs(img1,img2,pairs,[filename,'S']);
18
19 % LM optimization
20 H = LMfindH(pairs(bestIdx,:));
21 end
```

4.2.1 Match SURF Feature Function

```
1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi349@purdue.edu
4
5 function pairs=surfMatch(img1,img2,filename)
6 % This function is to perform SURF match process
7 I1 = rgb2gray(img1);
8 I2 = rgb2gray(img2);
9
10 % detect and generate SURF features
11 points1 = detectSURFFeatures(I1);
12 points2 = detectSURFFeatures(I2);
13
14 [features1,valid_points1] = extractFeatures(I1,points1);
15 [features2,valid_points2] = extractFeatures(I2,points2);
16
17 % match SURF features
18 indexPairs = matchFeatures(features1,features2);
19
20 matchedPoints1 = valid_points1(indexPairs(:,1),:);
21 matchedPoints2 = valid_points2(indexPairs(:,2),:);
22
23 pairs = [matchedPoints1.Location, matchedPoints2.Location];
24
25 % Plot the image
26 %pltPairs(img1,img2,valid_points1.Location,valid_points2.Location,pairs
27 %    (:,[2,1,4,3]),[filename,'SURF'])
27 pairs = pairs(:,[2,1,4,3]);
28
29 end
```

4.2.2 RANSAC

```
1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi349@purdue.edu
4
5 function [bestIdx,bestH] = ransacH(pairs)
6 % This function can find the most fitting Homography from pairs by RANSAC
7 % pairs is the set of all matched points.
8 nt= size(pairs,1); % Total number of matches
9 % parameters:
10 p = .99; % The probability that at least one of the N trials will be free of
11 % outliers.
11 m = 9; % the set of correspondences for calculating homography matrix H in each
11 trial.
12 esp = 0.2; % the rough estimation of false correspondences from the total set.
```

```

13 del = 20; % The decision threshold to construct the inlier set.
14 N = ceil(log(1-p)/log(1-(1-esp)^m))% Number of trials
15 M = floor((1-esp)*nt) % A minimum value for the size of the inlier set for it to be
   acceptable
16
17 Maxcount = 0;
18 bestH = zeros(3);
19 bestIdx = zeros(nt,1);
20 for i = 1:N
21     randRowIdx = randsample(nt,m);
22     H = findH(pairs(randRowIdx,:));
23     estPoints = H*[pairs(:,1:2)';ones(1,nt)];
24     estPoints = estPoints./ estPoints(end,:);
25     dist = sum(abs((estPoints(1:2,:) - pairs(:,3:4)').'));
26     inIdx = find(dist<del);
27     count = length(inIdx) % use a naive cheese board distance
28     if count > M && count > Maxcount
29         bestH = H
30         bestIdx = inIdx;
31         Maxcount = count
32     end
33 end
34 if sum(bestIdx) == 0
35     error('No inline pairs found!');
36 end
37
38
39 end

```

4.2.3 LSM Homography Calculator

```

1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi349@purdue.edu
4
5 function H = findH(pairs)
6 % This function can find homography by SVD
7 % pairs are m (m>5) pairs of corresponding points by (x1,y1,x2,y2)
8 m = size(pairs,1);
9 if m < 5
10    disp("No necessary number of pairs (m<5)");
11    return
12 end
13 disp([num2str(m)," pairs points are used to find the homography."]);
14 A = zeros(m*2,9);
15 % Construct A
16 for i = 0:m-1
17    A(i*2+1:i*2+2,:) = [0 0 0 -pairs(i+1,1) -pairs(i+1,2) -1 pairs(i+1,4)*pairs(i+1,1)
18                           pairs(i+1,4)*pairs(i+1,2) pairs(i+1,4);
19                           pairs(i+1,1) pairs(i+1,2) 1 0 0 0 -pairs(i+1,1)*pairs(i+1,3) -pairs(i+1,3)*
20                           pairs(i+1,2) -pairs(i+1,3)];
21 end
22 if rank(A)< 9
23    disp("Warning: Rank(A) is less than 9!");
24 end
25 [~,~,V] = svd(A); % V is order by the descending order of S
26 H = V(:,end); % H is the vector with smallest S
27 H = [H(1:3)';H(4:6)';H(7:9)'];
28 end

```

4.2.4 Levenberg-Marquardt Homography Calculator

```

1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi349@purdue.edu
4
5 function H = LMfindH(pairs)
6 pairs = double(pairs);
7 H0 = 1*ones(1,9);
8 fun = @(H) myfun(H, pairs);
9 options.Algorithm = 'levenberg-marquardt';
10 H = lsqnonlin(fun,H0,[],[],options);
11 H = [H(1:3);H(4:6);H(7:9)];

```

```

12
13 end
14
15 function err = myfun(H, pairs)
16 % This is the function for non linear least square optimzation
17 H = [H(1:3);H(4:6);H(7:9)];
18 n = size(pairs,1);
19 xdata = [pairs(:,1:2),ones(n,1)]';
20 ydata = [pairs(:,3:4),ones(n,1)]';
21 estydata = H*xdata;
22 estydata = estydata./estydata(end,:);
23 err = ydata - estydata;
24 end

```

4.2.5 SURF Matching Plot with Inliers and Outliers

```

1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi 349@purdue.edu
4
5 function pltPairs(C1,C2,pairs, pairsIdx, filename)
6 % This function is to generate the final image of SURF matching
7 % also indicates inliners and outliners
8
9 [h,w,c] = size(C1);
10 fig = figure;
11 imshow([C1,C2]);
12 hold on
13 V1 = pairs(:,[2,1]);
14 V2 = pairs(:,[4,3]);
15 plot(V1(:,1),V1(:,2), 'x');
16 hold on
17 plot(V2(:,1)+w,V2(:,2), 'o');
18 hold on
19 set(gca, 'position',[0 0 1 1], 'units', 'normalized')
20 x1 = pairs(:,1);
21 y1 = pairs(:,2);
22 x2 = pairs(:,3);
23 y2 = pairs(:,4)+w;
24
25 for i = 1:size(pairs,1)
26 if ismember(i, pairsIdx)
27 line([y1(i),y2(i)],[x1(i),x2(i)], 'Color', 'g', 'LineStyle', '-');
28 hold on
29 else
30 line([y1(i),y2(i)],[x1(i),x2(i)], 'Color', 'b', 'LineStyle', '-');
31 hold on
32 end
33 end
34 hold off;
35 saveas(fig,[filename, '.png'])
36 end

1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi349@purdue.edu
4
5 function H = LMfindH(pairs)
6 pairs = double(pairs);
7 H0 = 1*ones(1,9);
8 fun = @(H) myfun(H, pairs);
9 options.Algorithm = 'levenberg-marquardt';
10 H = lsqnonlin(fun,H0,[],[],options);
11 H = [H(1:3);H(4:6);H(7:9)];
12
13 end
14
15 function err = myfun(H, pairs)
16 % This is the function for non linear least square optimzation
17 H = [H(1:3);H(4:6);H(7:9)];
18 n = size(pairs,1);
19 xdata = [pairs(:,1:2),ones(n,1)]';
20 ydata = [pairs(:,3:4),ones(n,1)]';

```

```

21 estydata = H*xdata;
22 estydata = estydata./estydata(end,:);
23 err = ydata - estydata;
24 end

```

4.3 Projection Function by Homography

```

1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi349@purdue.edu
4
5 function img = projectH(img1,img2,H,mono,filename)
6 % This function can project img1 to img2 by H, x2 = H*x1
7 if nargin <4
8     filename = datetime('today');
9 end
10 % Mono switch
11 % mono = 0
12
13 [h1,w1,c1] = size(img1);
14 [h2,w2,c2] = size(img2);
15 % H = H'
16 % define new img size
17 points = [0 0 1; h1 0 1; 0 w1 1; h1 w1 1]';
18 newpoints = H*points;
19 newpoints = newpoints./newpoints(end,:);
20 minh1 = floor(min(newpoints(1,:)));
21 maxh1 = ceil(max(newpoints(1,:)));
22 minw1 = floor(min(newpoints(2,:)));
23 maxw1 = ceil(max(newpoints(2,:)));
24
25 minh = min([minh1,0]);
26 maxh = max([maxh1,h2]);
27 minw = min([minw1,0]);
28 maxw = max([maxw1,w2]);
29
30 newh = ceil(maxh - minh);
31 neww = ceil(maxw-minw);
32
33 img = zeros(newh,neww,c1);
34
35
36 for x = 1:h2
37     for y = 1:w2
38         if mono == 1
39             img(x-minh,y-minw,1) = img2(x,y,1);
40         else
41             img(x-minh,y-minw,:) = img2(x,y,:);
42         end
43     end
44 end
45 end
46
47 invH = H^-1;
48
49 for x = 1:newh
50     for y = 1:neww
51         Loc1 = invH*[x+minh,y+minw,1]';
52         Loc1 = round(Loc1./Loc1(end));
53         if Loc1(1) > 0 && Loc1(1) <= h1 && Loc1(2) > 0 && Loc1(2) <= w1
54             if mono == 1
55                 img(x,y,2) = img1(Loc1(1),Loc1(2),2);
56             else
57                 if sum(img1(Loc1(1),Loc1(2),:)) ~= 0
58                     img(x,y,:) = img1(Loc1(1),Loc1(2),:);
59                 end
60             end
61         end
62     end
63 end
64
65 if mono == 1
66     filename = [filename, 'M'];

```

```

67 end
68
69
70 img = uint8(img);
71 fig = figure;
72 set(gca, 'position',[0 0 1 1], 'units', 'normalized')
73 imshow(img)
74 saveas(fig,[filename, '.png'])
75 end

```

4.4 Mosaicing Function by Homography

```

1 %% ECE 661 2018 Fall Homework 5
2 % Ye Shi
3 % shi349@purdue.edu
4
5 function img = mosaicing5(imgset,H,filename)
6 % imgset is a 5-image cell
7 % H is a 4-pairwise-homography cell
8 if nargin <3
9     filename = datetime('today');
10 end
11 % Mono switch
12 mono = 1
13
14 % All images are the same size
15 [h1,w1,c1] = size(imgset{1});
16
17 % define new img size
18 points = [0 0 1; h1 0 1; 0 w1 1; h1 w1 1];
19 newpoints1 = H{1}*H{2}*points;
20 newpoints1 = newpoints1./newpoints1(end,:);
21 minh1 = floor(min(newpoints1(1,:)));
22 maxh1 = ceil(max(newpoints1(1,:)));
23 minw1 = floor(min(newpoints1(2,:)));
24 maxw1 = ceil(max(newpoints1(2,:)));
25
26 newpoints2 = H{2}*points;
27 newpoints2 = newpoints2./newpoints2(end,:);
28 minh2 = floor(min(newpoints2(1,:)));
29 maxh2 = ceil(max(newpoints2(1,:)));
30 minw2 = floor(min(newpoints2(2,:)));
31 maxw2 = ceil(max(newpoints2(2,:)));
32
33 newpoints3 = H{3}^-1 * points;
34 newpoints3 = newpoints3./newpoints3(end,:);
35 minh3 = floor(min(newpoints3(1,:)));
36 maxh3 = ceil(max(newpoints3(1,:)));
37 minw3 = floor(min(newpoints3(2,:)));
38 maxw3 = ceil(max(newpoints3(2,:)));
39
40 newpoints4 = H{3}^-1 * H{4}^-1 * points;
41 newpoints4 = newpoints4./newpoints4(end,:);
42 minh4 = floor(min(newpoints4(1,:)));
43 maxh4 = ceil(max(newpoints4(1,:)));
44 minw4 = floor(min(newpoints4(2,:)));
45 maxw4 = ceil(max(newpoints4(2,:)));
46
47 % Find the extrema of all projections
48 minh = min([minh1,minh2,minh3,minh4,0]);
49 maxh = max([maxh1,maxh2,maxh3,maxh4,h1]);
50 minw = min([minw1,minw2,minw3,minw4,0]);
51 maxw = max([maxw1,maxw2,maxw3,maxw4,w1]);
52
53 newh = ceil(maxh - minh);
54 neww = ceil(maxw-minw);
55
56 img = zeros(newh,neww,c1);
57
58 % middle image is the base
59 for x = 1:h1
60     for y = 1:w1

```

```

62     if mono == 1
63         img(x-minh,y-minw,1)= imgset{3}(x,y,1);
64     else
65         img(x-minh,y-minw,:)= imgset{3}(x,y,:);
66     end
67
68     end
69 end
70
71 invH = (H{1}*H{2})^-1;
72
73 % Left image
74 for x = 1:newh
75     for y = 1:neww
76         Loc1 = invH*[x+minh,y+minw,1]';
77         Loc1 = round(Loc1./Loc1(end));
78         if Loc1(1) > 0 && Loc1(1)<= h1 && Loc1(2) > 0 && Loc1(2)<= w1
79             if mono == 1
80                 img(x,y,2) = imgset{1}(Loc1(1),Loc1(2),2);
81             else
82                 if sum(imgset{1}(Loc1(1),Loc1(2),:)) ~= 0
83                     img(x,y,:)= imgset{1}(Loc1(1),Loc1(2),:);
84                 end
85             end
86         end
87     end
88 end
89
90
91 invH = H{2}^-1;
92 % Left-middle image
93 for x = 1:newh
94     for y = 1:neww
95         Loc1 = invH*[x+minh,y+minw,1]';
96         Loc1 = round(Loc1./Loc1(end));
97         if Loc1(1) > 0 && Loc1(1)<= h1 && Loc1(2) > 0 && Loc1(2)<= w1
98             if mono == 1
99                 img(x,y,3) = imgset{2}(Loc1(1),Loc1(2),3);
100            else
101                if sum(imgset{2}(Loc1(1),Loc1(2),:)) ~= 0
102                    img(x,y,:)= imgset{2}(Loc1(1),Loc1(2),:);
103                end
104            end
105        end
106    end
107 end
108
109
110
111 invH = H{3};
112 % Right-middle image
113 for x = 1:newh
114     for y = 1:neww
115         Loc1 = invH*[x+minh,y+minw,1]';
116         Loc1 = round(Loc1./Loc1(end));
117         if Loc1(1) > 0 && Loc1(1)<= h1 && Loc1(2) > 0 && Loc1(2)<= w1
118             if mono == 1
119                 img(x,y,3) = imgset{4}(Loc1(1),Loc1(2),3);
120             else
121                 if sum(imgset{4}(Loc1(1),Loc1(2),:)) ~= 0
122                     img(x,y,:)= imgset{4}(Loc1(1),Loc1(2),:);
123                 end
124             end
125         end
126     end
127 end
128
129 invH = H{4}*H{3};
130 % Right-middle image
131 for x = 1:newh
132     for y = 1:neww
133         Loc1 = invH*[x+minh,y+minw,1]';
134         Loc1 = round(Loc1./Loc1(end));

```

```

135 if Loc1(1) > 0 && Loc1(1)<= h1 && Loc1(2) > 0 && Loc1(2)<= w1
136     if mono == 1
137         img(x,y,2) = imgset{5}(Loc1(1),Loc1(2),2);
138     else
139         if sum(imgset{5}(Loc1(1),Loc1(2),:)) ~= 0
140             img(x,y,:)= imgset{5}(Loc1(1),Loc1(2),:);
141         end
142     end
143 end
144 end
145 end
146
147 if mono == 1
148     filename = [filename, 'M'];
149 end
150
151
152 img = uint8(img);
153 fig = figure;
154 set(gca, 'position',[0 0 1 1], 'units', 'normalized')
155 imshow(img)
156 saveas(fig,[filename, '.png'])
157 end

```