ENSE 885AY

Application of Deep Learning in Computer Vision

Assignment A03

Camera Calibration and Fundamental Matrix Estimation with RANSAC

Instructed by

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1. Introduction

1.1. Overview (Key points from the assignment description) [1]

Assignment Subject:

Camera Calibration and Fundamental Matrix Estimation with RANSAC

Assignment objectives:

- The goal of this project is to introduce you to camera and scene geometry
- Estimate the camera projection matrix, which maps 3D world coordinates to 2D image coordinates.
- Estimate the fundamental matrix, which relates points in one scene to epipolar lines in another perspective of the same scene.
- Estimating the fundamental matrix for the correspondences of two images using the RANSAC model-fitting algorithm.

Steps to local feature matching between two images (image 1 & image 2):

- 1. Estimating the projection matrix:

 - ⇒ calculate_camera_center()
- 2. Estimating the fundamental matrix:
 - ⇒ estimate_fundamental matrix()
- 3. Estimating the fundamental matrix with unreliable ORB matches using RANSAC:
 - ransac_fundamental_matrix()

2. Student Code

2.1. Estimating the projection matrix (calculate_projection_matrix() & calculate_camera_center())

Algorithm and implementation of calculate_projection_matrix() [1]:

The first step is to set up a system of equations using the corresponding 2D and 3D points:

$$\begin{pmatrix} u \\ v \\ 1 \end{pmatrix} \cong \begin{pmatrix} u * s \\ v * s \\ s \end{pmatrix} = \begin{pmatrix} m_{11} & m_{12} & m_{13} & m_{14} \\ m_{21} & m_{22} & m_{23} & m_{24} \\ m_{31} & m_{32} & m_{33} & m_{34} \end{pmatrix} \begin{pmatrix} X \\ Y \\ Z \\ 1 \end{pmatrix}$$

$$u = \frac{m_{11}X + m_{12}Y + m_{13}Z + m_{14}}{m_{31}X + m_{32}Y + m_{33}Z + m_{34}}$$

$$\to (m_{31}X + m_{32}Y + m_{33}Z + m_{34})u = m_{11}X + m_{12}Y + m_{13}Z + m_{14}$$

$$\to 0 = m_{11}X + m_{12}Y + m_{13}Z + m_{14} - m_{31}uX - m_{32}uY - m_{33}uZ - m_{34}u$$

$$v = \frac{m_{21}X + m_{22}Y + m_{23}Z + m_{24}}{m_{31}X + m_{32}Y + m_{33}Z + m_{34}}$$

$$\to (m_{31}X + m_{32}Y + m_{33}Z + m_{34})v = m_{21}X + m_{22}Y + m_{23}Z + m_{24}$$

$$\to 0 = m_{21}X + m_{22}Y + m_{23}Z + m_{24} - m_{31}vX - m_{32}vY - m_{33}vZ - m_{34}v$$

$$\to 0 = m_{21}X + m_{22}Y + m_{23}Z + m_{24} - m_{31}vX - m_{32}vY - m_{33}vZ - m_{34}v$$

The above system can be transformed to following matrix form:

$$A_{2N\times 11} * M_{11\times 1} = b_{2N\times 1}$$

There are two different rows in matrices A and b (row[2k] and row[2k+1]:

$$\begin{bmatrix} row_{2k}^A \\ row_{2k+1}^A \end{bmatrix}_{2N\times 11} M_{11\times 1} = \begin{bmatrix} row_{2k}^b \\ row_{2k+1}^b \end{bmatrix}_{2N\times 1}$$

where

$$\begin{bmatrix} row_{2k}^A \\ row_{2k+1}^A \end{bmatrix}_{2\times 11} = \begin{bmatrix} X_k & Y_k & Z_k & 1 & 0 & 0 & 0 & -u_k X_k & -u_k Y_k & -u_k Z_k \\ 0 & 0 & 0 & X_k & Y_k & Z_k & 1 & -v_k X_k & -v_k Y_k & -v_k Z_k \end{bmatrix}_{2\times 11}$$

$$\begin{bmatrix} row_{2k}^b \\ row_{2k+1}^b \end{bmatrix}_{2\times 1} = \begin{bmatrix} u_k \\ v_k \end{bmatrix}_{2\times 1}$$

Matrices A and b will be constructed using the corresponding 2D and 3D points as below:

Define constants and placeholders for matrices A and b

```
# Matrices A and b will be constructed using the corresponding 2D and 3D points as below
# Define constants and placeholders for matrices A and b
N = int(points_2d.shape[0])
b = np.zeros((2*N,1))
A = np.zeros((2*N,11))
```

For every row[k] in the corresponding 2D and 3D points

```
Construct row[2k] and row[2k+1] of matrices A and b
```

```
# For every row[k] in the corresponding 2D and 3D points
for k in range(0,N):
    row1 = 2*k
    row2 = 2*k+1

# Construct row[2k] and row[2k+1] of matrices A and b
    b[row1,0] = points_2d[k,0]
    b[row2,0] = points_2d[k,1]

A[row1,0:3] = points_3d[k,:]
    A[row1,3] = 1
    A[row1,8:11] = -points_2d[k,0]*points_3d[k,:]

A[row2,4:7] = points_3d[k,:]
    A[row2,7] = 1
    A[row2,8:11] = -points_2d[k,1]*points_3d[k,:]
```

Solve A* M_{11x1} =b using np.linalg.lstsq() to obtain M_{11x1} # Solve A*M 11 = b using np.linalg.lstsq() to obtain M 11

```
M11 = np.linalg.lstsq(A, b, rcond = None)[0]
```

```
Append M_{34}=1 to M_{11x1} and reshape it as M_{3x4} # Append M_34 = 1 to M_11 and reshape it as M_3x4 M12 = np.append(M11, [1]) M = M12.reshape((3, 4))
```

Algorithm and implementation of calculate_camera_center() [1]:

Camera center is calculated using following equations:

Extract matrices Q and m4 from matrix M

```
\begin{split} M &= [M_{3\times3}|M_{3\times1}];\\ Q &= M_{3\times3}\\ m_4 &= M_{3\times1}\\ &\quad \text{\# Extract matrices Q and m4 from matrix M}\\ Q &= \text{M[0:3,0:3]}\\ &\quad \text{m4} &= \text{M[:,3]} \end{split}
```

Calculate camera center (matrix cc)

```
cc = -Q^{-1}m_4 # Calculate camera center (matrix cc) cc = -np.linalg.inv(Q) @ m4
```

2.2. Estimating the fundamental matrix (estimate_fundamental matrix())

Algorithm and implementation of estimate_fundamental matrix() [1]:

The first step is to set up a system of equations using the corresponding 2D points a and b:

$$\begin{pmatrix} u' & v' & 1 \end{pmatrix} \begin{pmatrix} f_{11} & f_{12} & f_{13} \\ f_{21} & f_{22} & f_{23} \\ f_{31} & f_{32} & f_{33} \end{pmatrix} \begin{pmatrix} u \\ v \\ 1 \end{pmatrix} = 0$$

$$(u' \quad v' \quad 1) \begin{pmatrix} f_{11}u + f_{12}v + f_{13} \\ f_{21}u + f_{22}v + f_{23} \\ f_{31}u + f_{32}v + f_{33} \end{pmatrix} = 0$$

$$(f_{11}uu' + f_{12}vu' + f_{13}u' + f_{21}uv' + f_{22}vv' + f_{23}v' + f_{31}u + f_{32}v + f_{33}) = 0$$

The above equation can be transformed to following matrix form:

$$UV_{N\times9} * F_{9\times1} = 0_{N\times1}$$

$$[[u_k^A u_k^B]_{N \times 1} \quad [v_k^A u_k^B]_{N \times 1} \quad [u_k^B]_{N \times 1} \quad [u_k^A v_k^B]_{N \times 1} \quad [u_k^A v_k^B]_{N \times 1} \quad [v_k^B]_{N \times 1} \quad [u_k^A]_{N \times 1} \quad [u_k^A]_{N \times 1} \quad [u_k^A]_{N \times 1} \quad [u_k^A v_k^B]_{N \times 1} \quad [u_k^A v_$$

$$= \begin{bmatrix} 0 \\ \vdots \\ 0 \end{bmatrix}_{N \times 1}$$

Matrix UV will be constructed using the corresponding 2D points_a and points_b as below:

Set with_normalization to True or False

```
# Set with_normalization to True or False
with_normalization = True
```

If with_normalization = True: Normalize points_a and points_b and obtain their transformation matrices T_a and T_b

```
# If with_normalization = True:
# Normalize points_a and points_b and obtain T_a and T_b
if with_normalization == True:
    points_a, T_a = normalization_function(points_a)
    points_b, T_b = normalization_function(points_b)
```

Define constant N, and extract uA, vA, uB, vB, and I_N from points_a and points_

Construct matrix UV using uA, vA, uB, vB, and I_N and their multiplications

```
# Extract uA, vA, uB, vB, and I_N from points_a and points_b
uA = points_a[:,0].reshape(N,1)
vA = points_a[:,1].reshape(N,1)
uB = points_b[:,0].reshape(N,1)
vB = points_b[:,1].reshape(N,1)
I_N = np.ones((N,1))

# Construct matrix UV using uA, vA, uB, vB, and I_N and their multiplications
UV = np.hstack((uA*uB, vA*uB, uB, uA*vB, vA*vB, vB, uA, vA, I_N ))
```

```
Solve UV* F_{9x1} = 0 using np.linalg. svd () to obtain full-rank F_{9x1} # Solve UV* F_{9x1} = 0 to obtain full-rank F_{9x1} _, _, vh0 = np.linalg.svd(UV,full_matrices = True) F3 = vh0[8,:].reshape(3,3)
```

Reduce F rank from 3 to 2

```
# Reduce F rank from 3 to 2
u3, s3, vh3 = np.linalg.svd(F3, full_matrices = True)
s3[2] = 0
s2 = np.diag(s3)
F = np.dot(u3, np.dot(s2, vh3))
```

If with_normalization = True: Transform F_norm to F_orig

Algorithm and implementation of normalization_function() [1]:

Define constant N

```
# Define constant N
N = points_a.shape[0]
```

Obtain center of points

```
center_a = [\overline{u_a} \quad \overline{v_a}] # Obtain center of points center_a = np.average(points_a,axis = 0)
```

Center points around the center

```
centered\_a_{N\times 2} = \begin{bmatrix} u_a & v_a \end{bmatrix}_{N\times 2} - \begin{bmatrix} \overline{u_a} & \overline{v_a} \end{bmatrix} = \begin{bmatrix} u_a - \overline{u_a} & v_a - \overline{v_a} \end{bmatrix}_{N\times 2} # Center points around the center centered_a = points_a-center_a.reshape(1,2)
```

Obtain standard deviation

$$std_a = \sqrt{\frac{([u_a - \overline{u_a} \quad v_a - \overline{v_a}]_{N \times 2})^2}{2N}}$$
 # Obtain standard deviation
$$std_a = \text{np.sqrt((np.sum((centered_a)**2,axis = None)/(2*N)))}$$

Calculate scale

$$scale_a = 1/std_a$$
 # Calculate scale $scale_a = 1/std_a$

Obtain scaled and centered points_a

```
scaled-centered_a = scale_a*centered\_a_{N\times 2}  \begin{tabular}{l} \# \ Obtain \ scaled \ and \ centered \ points\_a \ scaled\_centered\_a = scale\_a*centered\_a \end{tabular}
```

Obtain Ts and Tc matrices

$$Ts_{a} = \begin{bmatrix} scale_{a} & 0 & 0 \\ 0 & scale_{a} & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

$$Tc_{a} = \begin{bmatrix} 1 & 0 & -\overline{u}_{a} \\ 0 & 1 & \overline{v}_{a} \\ 0 & 0 & 1 \end{bmatrix}$$

```
# Obtain Ts and Tc matrices
Ts_a = np.diag(np.array([scale_a, scale_a, 1]))
Tc_a = np.diag(np.array([1., 1., 1.]))
Tc_a[0,2] = -center_a[0]
Tc a[1,2] = -center a[1]
```

Calculate matrix T

```
T_a = Ts_a * Tc_a # Calculate matrix T T_a = Ts_a @ Tc_a
```

2.3. Estimating the fundamental matrix with unreliable ORB matches using RANSAC (ransac_fundamental_matrix())

Algorithm and implementation of ransac_fundamental_matrix() [1]:

Define constants

```
# Define constants
N = matches_a.shape[0]
max_iter = 15000
threshold = 0.02
batch_size = 8
```

Define random indexes with

```
Range = 0 : N = number of points

Size = max_iter x batch_size = 15000 x 8

# Define random indexes with size = (max_iter, batch_size)
rand_idx = np.random.randint(N,size = (max_iter, batch_size))
```

Obtain UV matrix (algorithm described in previous section)

```
# Obtain UV matrix (algorithm described in previous section)
# Extract uA, vA, uB, vB, and I_N from points_a and points_b
uA = matches_a[:,0].reshape(N,1)
vA = matches_a[:,1].reshape(N,1)
uB = matches_b[:,0].reshape(N,1)
vB = matches_b[:,1].reshape(N,1)
I_N = np.ones((N,1))
# Construct matrix UV using uA, vA, uB, vB, and I_N and their multiplications
UV = np.hstack((uA*uB, vA*uB, uB, uA*vB, vA*vB, vB, uA, vA, I_N ))
```

Define placeholder for inlier_count

```
# Define placeholders
inlier_count = np.zeros(max_iter)
```

Iteration for k=0:max_iter

Estimate F matrix using random batch[k] of 8 pairs of points

Calculate cost of estimated F using following equation

$$cost = abs(UV_{N\times9} * F_{9\times1} - 0_{N\times1})$$

```
cost = abs([[u_{k}^{A}u_{k}^{B}]_{N\times 1} \quad [v_{k}^{A}u_{k}^{B}]_{N\times 1} \quad [u_{k}^{B}]_{N\times 1} \quad [u_{k}^{A}v_{k}^{B}]_{N\times 1} \quad [u_{k}^{A}v_{k}^{B}]_{N\times 1} \quad [v_{k}^{B}]_{N\times 1} \quad [u_{k}^{A}]_{N\times 1} \quad [
```

```
\begin{vmatrix}
f_{12} \\
f_{13} \\
f_{21} \\
* & f_{22} \\
f_{23} \\
f_{31} \\
f_{32} \\
f_{24}
\end{vmatrix}
```

Obtain and save number of inliers (with cost < threshold)

```
# Iteration for k=0:max_iter
```

```
for k in range(max_iter):
        # Estimate F matrix using random batch[k] of 8 pairs of points
        F = estimate_fundamental_matrix(matches_a[rand_idx[k,:],:],matches_b[rand_idx[k,:],:]
)
        # Calculate cost of estimated F using following equation
        cost k = np.abs(UV @ F.reshape((9,1)))
        # Obtain and save number of inliers (with cost < threshold)
        inlier idx = cost k < threshold</pre>
        inlier_count[k] = np.sum(inlier_idx)
Sort inlier_count from maximum to minimum
    # Sort inlier count from maximum to minimum
    sort_idx = np.argsort(-inlier_count)
Obtain best_batch with maximum inliers
    # Obtain best batch with maximum inliers
    best_idx = sort_idx[0]
    best_batch = rand_idx[best_idx,:]
Estimate best_F matrix using best_batch
    # Estimate best F matrix using best batch
    best_F = estimate_fundamental_matrix(matches_a[best_batch,:], matches_b[best_batch,:])
Calculate best_cost of estimated best_F
    # Calculate best cost of estimated best F
    best_cost = np.abs( UV @ best_F.reshape((9,1)) )
Obtain and save number of inliers (with best_cost < threshold)
    # Obtain and save number of inliers (with best cost < threshold)
    inlier idx = best cost < threshold
    best_inlier_count = np.sum(inlier_idx)
```

Sort cost from minimum (best match) to maximum (worst match)

```
# Sort cost from minimum (best match) to maximum (worst match)
index=np.argsort(best_cost[:,0])[:50]
```

Obtain best pairs of matching points with minimum cost

```
# Obtain best pairs of matching points with minimum cost
inliers_a=matches_a[index,:]
inliers_b=matches_b[index,:]
```

Print results

```
# Print results
print('Found', best_inlier_count,'inliers / ', N, 'points')
print('inliers / total points :', int(100*best_inlier_count/N), '%')
```

3. Results and Discussion

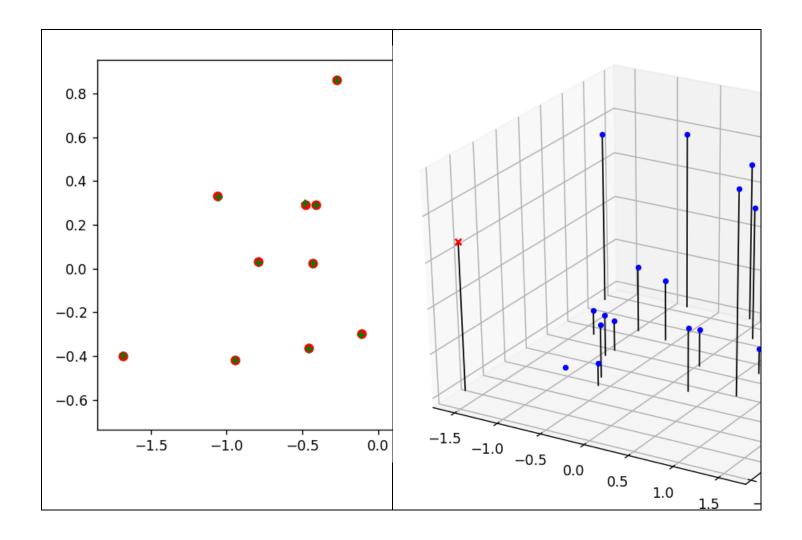
3.1. Estimation of the projection matrix and camera center

Table 1: Estimation of the projection matrix and camera center

Projection Matrix	Total Residual	Camera Center
[0.76785834 -0.49384797 -0.02339781 0.00674445]	0.044535	<-1.5126, -2.3517, 0.2827>
[-0.0852134 -0.09146818 -0.90652332 - 0.08775678]		
[0.18265016 0.29882917 -0.07419242 1.]		

Table 2: Point Visualization and Point 3D View

Point Visualization	Point 3D View

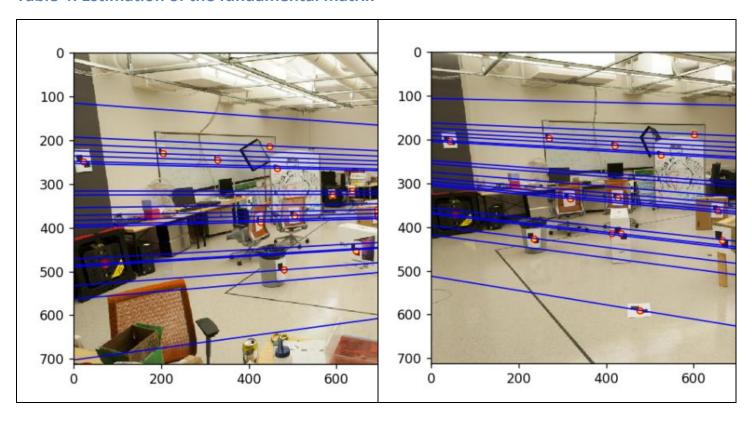


3.2. Estimation of the fundamental matrix

Table 3: Estimation of the fundamental matrix

Fundamental Matrix without Normalization	Fundamental Matrix with Normalization
([[-5.36264198e-07, 7.90364771e-06, - 1.88600204e-03],	([[-1.17248591e-07, 1.60824663e-06, - 4.01980786e-04],
[8.83539184e-06, 1.21321685e-06, 1.72332901e-02],	[1.11212887e-06, -2.73443755e-07, 3.23319884e-03],
[-9.07382264e-04, -2.64234650e-02, 9.99500092e-01]])	[-2.36400817e-05, -4.44404958e-03, 1.03455561e-01]])

Table 4: Estimation of the fundamental matrix



3.3. Estimation of the fundamental matrix with unreliable ORB matches using RANSAC

For four pairs of images, fundamental matrix was estimated with / without normalization. The results are summarized in following table:

For all images expect Woofruff, normalization has a positive effect on number of inliers.

Table 5: Inlier count for fundamental estimated with / without normalization

		inliers / total points	inliers / total points %
Notre Dame	Without normalization	626 / 1282	48 %
	With normalization	1043 / 1282	81 %

Mount Rushmore	Without normalization	519 / 1177	44 %
	With normalization	547 / 1177	46 %
Gaudi	Without normalization	350 / 1037	33 %
	With normalization	478 / 1037	46 %
Woodruff	Without normalization	506 / 1137	44 %
	With normalization	424 / 1137	37 %

Table 6: Result for Notre Dame without Normalization

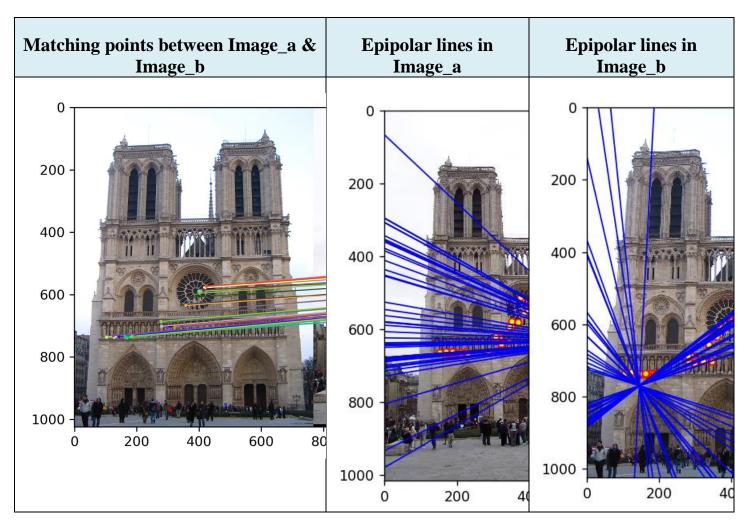


Table 7: Result for Notre Dame with Normalization

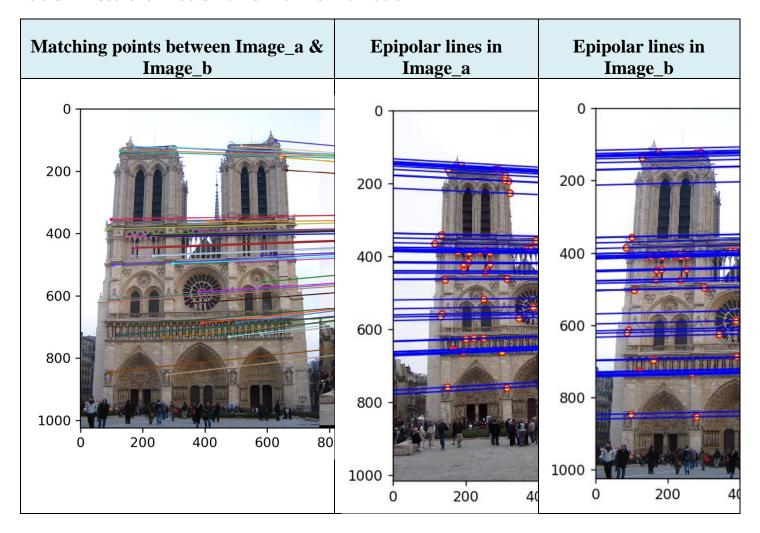


Table 8: Results for Mount Rushmore without Normalization

Matching points between Image_a & Image_b	
---	--

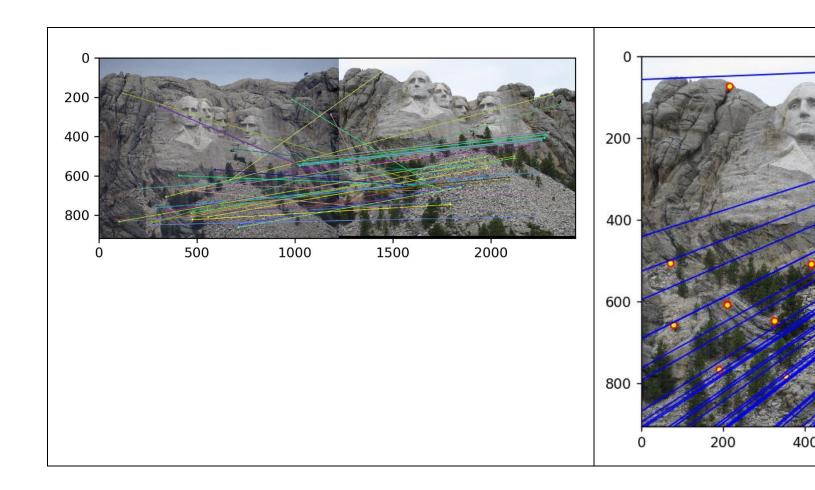


Table 9: Results for Mount Rushmore with Normalization

Matching points between Image_a & Image_b	Epipolar lines in Image_a	Epipolar lines in Image_b
--	---------------------------	---------------------------

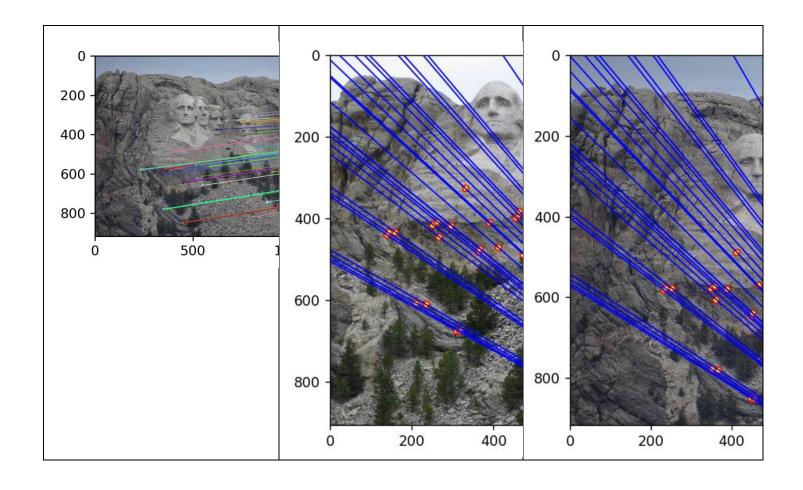


Table 10: Results for Gaudi without Normalization

Matching points between Image_a & Image_b	Epipolar lines in Image_a	Epipolar lines in Image_b
--	---------------------------	---------------------------

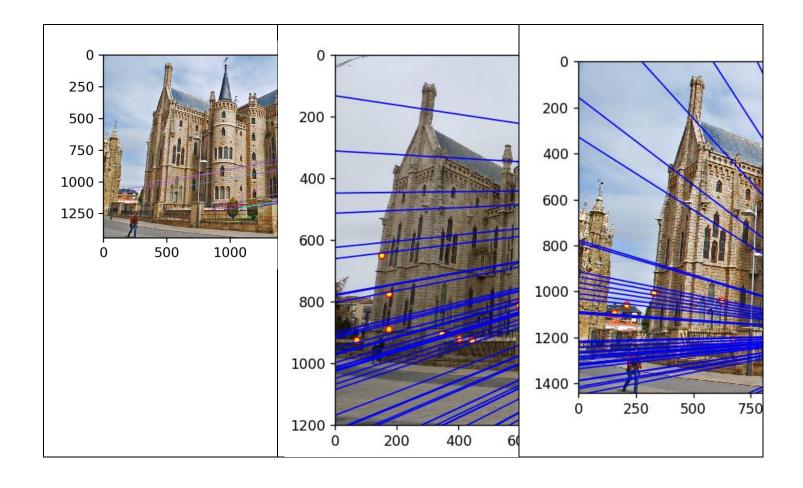


Table 11: Results for Gaudi with Normalization

Matching points between Image_a & Image_b	Epipolar lines in Image_a	Epipolar lines in Image_b
--	---------------------------	---------------------------

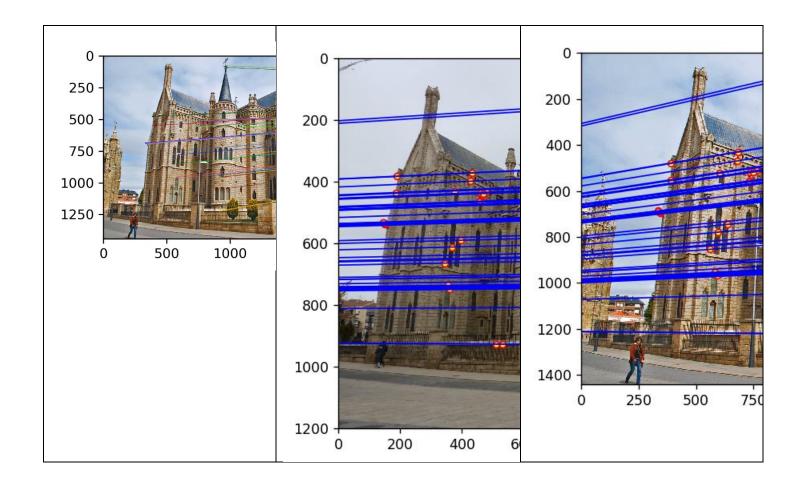


Table 12: Results for Woodruff without Normalization

	Matching points between Image_a & Image_b	Epipolar lines in Image_a	Epipolar lines in Image_b	
--	--	---------------------------	---------------------------	--

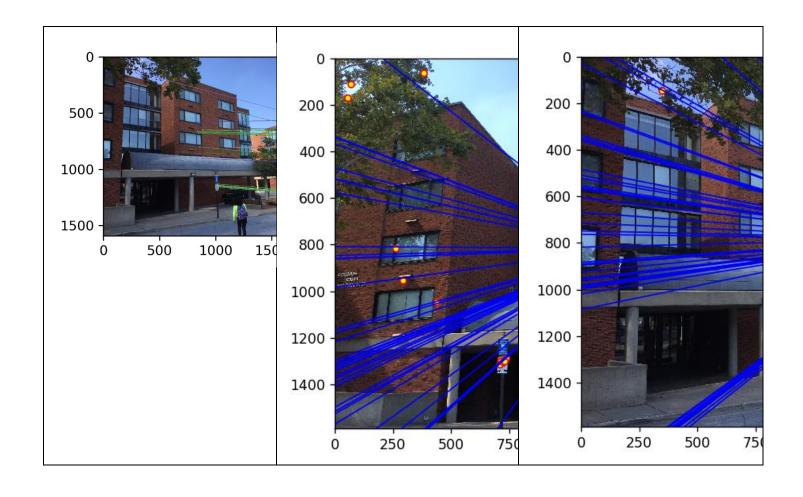
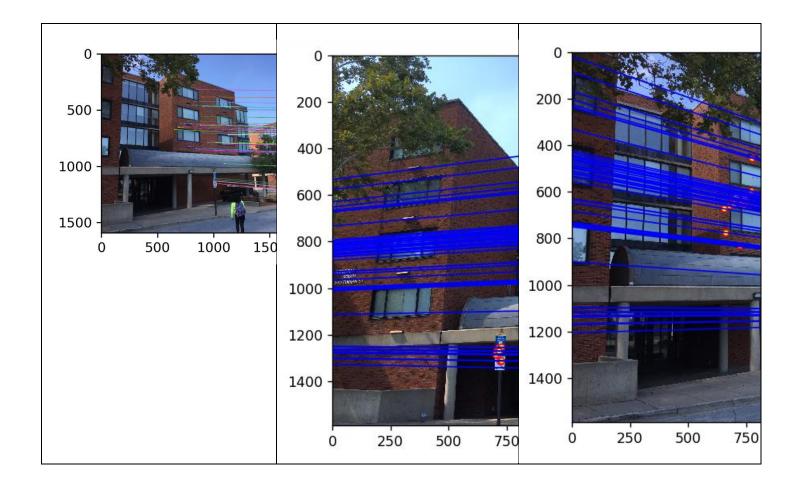


 Table 13: Results for Woodruff with Normalization

Matching points between Image_a & Image_b	Epipolar lines in Image_a	Epipolar lines in Image_b
--	---------------------------	---------------------------



3.4. Tuning threshold for ransac_fundamental_matrix()

I have written and used this function to obtain best threshold for ransac_fundamental_matrix().

Algorithm and implementation of tune_ransac_fundamental_matrix() [1]:

The general algorithm is the same as ransac_fundamental_matrix() with following differences:

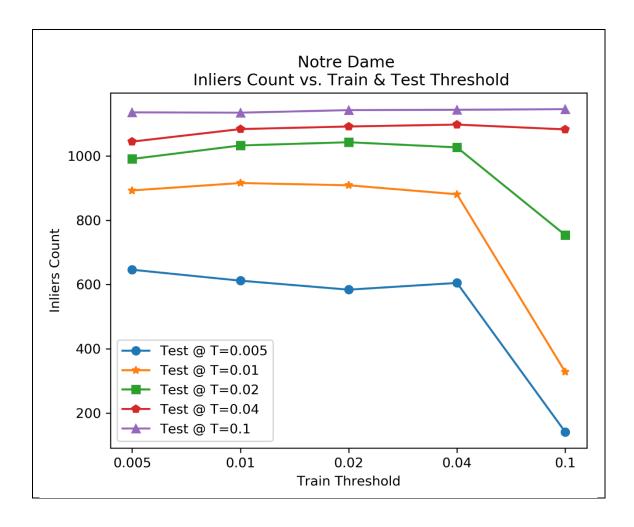
- Instead of a single threshold value, 5 different values are used:
 - $\circ \ threshold = [0.005, 0.01, 0.02, 0.04, 0.1]$
- I have also tried different threshold for training and testing:
 - Train threshold is the threshold used for first loop (that obtains best random batch of points);
 - Test threshold is the threshold used for best_F;
- best_inlier_count[i,j] is obtained for every train_thresh_j and test_thresh_i

```
o train thresh j = threshold[j]
        o test_thresh_i = threshold[i]
def tune ransac fundamental matrix(matches a, matches b):
   threshold = [0.005, 0.01, 0.02, 0.04, 0.1]
   For every train thresh j and test thresh i
       Obtain and save best inlier count[i,j]
   .....
   # TODO: YOUR RANSAC CODE HERE
   # Define constants
   N = matches a.shape[0]
   max iter = 15000
   threshold = [0.005, 0.01, 0.02, 0.04, 0.1]
   batch size = 8
   # Define random indexes with size = (max iter, batch size)
   rand_idx = np.random.randint(N,size = (max_iter, batch_size))
   # np.save('rand_idx_.npy',rand_idx)
   # rand idx = np.load('rand idx 526.npy')
   # Obtain UV matrix (algorithm described in previous section)
   # Extract uA, vA, uB, vB, and I N from points a and points b
   uA = matches a[:,0].reshape(N,1)
   vA = matches a[:,1].reshape(N,1)
   uB = matches_b[:,0].reshape(N,1)
   vB = matches_b[:,1].reshape(N,1)
   I_N = np.ones((N,1))
   # Construct matrix UV using uA, vA, uB, vB, and I N and their multiplications
   UV = np.hstack((uA*uB, vA*uB, uB, uA*vB, vA*vB, vB, uA, vA, I_N))
   # Define placeholders
   inlier count = np.zeros((len(threshold), max iter))
   best idx = np.zeros(len(threshold)).astype(int)
   best_inlier_count = np.zeros((len(threshold),len(threshold))).astype(int)
   # Iteration for k=0:max iter
   for k in range(max_iter):
       # Estimate F matrix using random batch[k] of 8 pairs of points
```

```
F = estimate_fundamental_matrix(matches_a[rand_idx[k,:],:],matches_b[rand_idx[k,:],:]
)
       # Calculate cost of estimated F using following equation
       cost_k = np.abs(UV @ F.reshape((9,1)))
       # Obtain and save number of inliers (with cost < threshold j)</pre>
       for j, train_thresh_j in enumerate(threshold):
          inlier_idx = cost_k < train_thresh_j</pre>
          inlier_count[j,k] = np.sum(inlier_idx)
   # For every train_thresh_j and test_thresh_i
   # Obtain and save best_inlier_count[i,j]
   for j, train_thresh_j in enumerate(threshold):
       sort_idx = np.argsort(-inlier_count[j,:])
       best_idx[j] = sort_idx[0]
       best_batch = rand_idx[best_idx[j],:]
       best_F = estimate_fundamental_matrix(matches_a[best_batch,:], matches_b[best_batch,:]
)
      cost = np.abs( UV @ best_F.reshape((9,1)) )
       for i, test thresh i in enumerate(threshold):
          inlier_idx = cost < test_thresh_i</pre>
          best_inlier_count[i,j] = np.sum(inlier_idx)
   # END OF YOUR CODE
   return best_inlier_count
```

Table 14: Inliers Count vs. Train & Test Threshold for Notre Dame

Inliers Count vs. Train & Test Threshold for Notre Dame



Remarks on the Inliers Count vs. Train & Test Threshold:

- Obviously, the higher test threshold value, the higher number of inliers. Therefore, it is not reasonable to tune threshold value using same value for training and testing;
- Also, higher inliers is not necessarily desirable because it increases the chance of false inliers;
- Therefore, it is best to compare different train thresholds based on constant test threshold;
- For test threshold = 0.02, highest inlier is obtained for train threshold = 0.02;
- Therefore, threshold value was chosen to be 0.02;

Extra Works

Following tasks were done outside assignment requirements:

Tuning threshold for ransac_fundamental_matrix()

References

- [1] Assignment 01 description by Dr. Kin-Choong Yow
- [2] Szeliski, R. (2010). Computer vision: algorithms and applications. Springer Science & Business Media.