## LAB ASSIGNMENT 7

ETH ZURICH, COMPUTER SCIENCE DEPARTMENT

# Computer vision

# Report

# Structure from Motion and RANSAC

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The objective of this lab assignment was to implement a Structure from Motion pipeline and RANSAC line fitting.

### 1 Structure from Motion

#### 1.1 Essential matrix estimation

The first step is to lift the keypoints to the normalized image plane first – I make the points homogeneous by adding ones as the last coordinates, calculate the inverse matrix of K and multiply it by homogeneous keypoints (transposed to match the dimensions of K matrix for multiplication, then I transpose normalized keypoints back to their initial dimensions). The next step is to create a constraint matrix to reformulate the constraint  $\hat{x}_1^T E \hat{x}_2 = 0$ . I do that by adding rows corresponding to each keypoint match in a form of  $[x_2x_1, x_2y_1, x_2z_1, y_2x_1, y_2y_1, y_2z_1, x_1z_2, y_1z_2, z_1z_2]$ , where in fact Z coordinates should be equal to 1. Then, the Singular Value Decomposition is applied to the constraint matrix and the last row of obtained  $V^H$  matrix contains elements of  $\hat{E}$ . This vector has to be reshaped to be a matrix  $3x_3$ . We need to fulfill the essential matrix constraints that the third singular value is equal to 0 and the first to are equal and greater than 0 (in our case we can assume them as 1). Therefore, SVD needs to be applied to  $\hat{E}$  and obtained U and  $V^H$  elements are multiplied by S, which is a diagonal matrix with 1, 1, 0 on its diagonal.

The code implementation of this part is presented below:

```
1 def EstimateEssentialMatrix(K, im1, im2, matches):
2
    K \text{ inv} = np.linalg.inv(K)
3
4
     kpts1 = MakeHomogeneous(im1.kps, ax=1)
5
     kpts2 = MakeHomogeneous(im2.kps, ax=1)
6
     normalized kps1 = K inv @ kpts1.T
7
     normalized kps2 = K inv @ kpts2.T
8
     normalized_kps1 = normalized_kps1.T
9
     normalized kps2 = normalized kps2.T
10
11
12
     constraint matrix = np.zeros((matches.shape[0], 9))
     for i in range(matches.shape[0]):
13
       x1,y1,z1 = normalized_kps1[i][0], normalized_kps1[i][1], normalized_kps1[i][2]
14
       x^2, y^2, z^2 = normalized kps^2[i][0], normalized kps^2[i][1], normalized kps^2[i][2]
15
16
       A_i = np. array([x2*x1, x2*y1, x2*z1, y2*x1, y2*y1, y2*z1, x1*z2, y1*z2, z1*z2])
17
       constraint_matrix[i] = A_i
18
     _, _, vh = np.linalg.svd(constraint_matrix)
19
     vectorized E hat = vh[-1,:]
20
21
    E hat = vectorized E hat.reshape((3,3))
22
23
    U\_hat\,,\ \_,\ VH\_hat\,=\ np\,.\,lin\,a\,l\,g\,.\,svd\,(\,E\ hat\,)
24
    s = np.array([1, 1, 0])
25
    S hat = np.diag(s)
26
    E = U hat @ S hat @ VH hat
27
28
29
     return E
```

#### 1.2 Next steps

I did not manage to overcome some of the problems and therefore did not continue on Structure from Motion task.

### 2 RANSAC

#### 2.1 Least-squares solution

To implement the least square solution, I first (as suggested by the linal lastsq documentation) rewrite the problem from y = kx + b to y = Ap, where A = [[x1]] and p = [[k], [b]]. Then I use the last to solve for p as presented in the code:

```
1 def least_square(x,y):

2 A = np.vstack([x, np.ones(len(x))]).T

3 k, b = np.linalg.lstsq(A, y, rcond=None)[0]

4 return k, b
```

#### 2.2 RANSAC implementation

First I use the np.choice function to randomly choose a subset of both x and y point sets. Then I compute the least-squares solution for this subset and with obtained k, b I calculate the number of inliers and the mask that denotes the indices of inliers (a point is an inlier if its distance to the line smaller than thres\_dist, which is compared in num\_inlier function). If the number of inliers is larger than the current best result, I update the parameters k\_ransac, b\_ransac, best\_inliers and inlier\_mask. The code corresponding to this part can be found below:

```
def num inlier(x,y,k,b,n samples,thres dist):
1
2
       num = 0
3
       mask = np.zeros(x.shape, dtype=bool)
4
       for i in range(n samples):
5
            dist = abs(k*x[i] - y[i] + b)/math.sqrt(k*k + 1)
6
            if dist < three dist:
7
                \operatorname{num} += 1
8
                mask[i] = True
9
10
       return num, mask
11
12
  def ransac(x,y, iter, n samples, thres dist, num subset):
       k \text{ ransac} = None
13
14
       b ransac = None
15
       inlier mask = None
       best inliers = 0
16
17
       for _ in range(iter):
18
           x\_subset = np.random.choice(x, size=num\_subset)
19
           y\_subset = np.random.choice(y, size=num subset)
20
21
           k, b = least square(x subset, y subset)
22
           num_inl, mask = num_inlier(x, y, k, b, n_samples, thres_dist)
            if num inl > best inliers:
23
24
                best inliers = num inl
25
                inlier mask = mask
26
                k \text{ ransac} = k
27
                b ransac = b
28
29
       return k ransac, b ransac, inlier mask
```

### 2.3 Results

After running RANSAC, the results presented in Fig.1 are obtained. The plot corresonds to:

- true coefficients: k = 1, b = 10,
- linear regression coefficients: k = 0.615965, b = 8.961727,
- RANSAC coefficients: k = 0.936744, b = 9.860836.



Figure 1: The result of running RANSAC for line fitting.