# **EECS 442 HW3**

# #1 - RANSAC

## 1.1

1) We need 2 points to compute a putative model. The formula for deriving the slope explains this - (y2 - y1) / (x2 - x1).

2)  

$$(1 - r)^{S} =$$
  
 $(1 - 0.1)^{2} =$   
 $(0.99)^{2} = 0.98$ 

3)  

$$P_{\text{no-outlier}} = (1 - (1 - r)^{S})^{N}$$

$$.95 = (1 - (1 - r)^{S})^{N}$$

$$.95 = (1 - (0.99)^{2})^{N}$$

$$.95 = (1 - 0.9801)^{N}$$

$$In(.95/0.199) = N$$

$$N = 1.56 \rightarrow 2 \text{ trials}$$

## 1.2

1) M has 4 degrees of freedom & 2 samples are required to find M

# 2) Missing

|          | find an M such that: M = arguin &   M= - y;  |
|----------|--|
|          | $M = \begin{bmatrix} m_{11} & m_{12} \\ m_{21} & m_{23} \end{bmatrix}$   |
| M(2) - 3 | $ \frac{y_{i1}}{m_{2i}} = \begin{bmatrix} m_{11} & m_{12} \\ m_{2i} & m_{2i} \end{bmatrix} \begin{bmatrix} x_{id} \\ x_{i2} \end{bmatrix} - \begin{bmatrix} y_{i1} \\ y_{i2} \end{bmatrix} $ $ = \begin{bmatrix} m_{11}x_{i1} + m_{12}x_{i2} \\ y_{i2} \end{bmatrix} - \begin{bmatrix} y_{i1} \\ y_{i3} \end{bmatrix} $  |
|          | $ \begin{array}{cccccccccccccccccccccccccccccccccccc$  |
|          | $ \begin{array}{cccccccccccccccccccccccccccccccccccc$  |
| M=; - ;  | $ \frac{1}{y_{i}} = a_{i}m - y_{i} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} =   [a_{i}]  ^{2} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} =   [a_{i}]  ^{2} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} $ $   M_{x_{i}} - y_{i}  ^{2} =   a_{i}m - y_{i}  ^{2} $   |
| A = [ A  | $\begin{bmatrix} \chi_{i_1} & \chi_{i_2} & 0 & 0 \\ 0 & 0 & \chi_{i_1} & \chi_{i_2} \\ \vdots & \ddots & \ddots & \vdots \\ \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \vdots \\ \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \vdots \\ \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \vdots \\ \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \vdots \\ \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \vdots \\ \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \chi_{i_n} & \vdots \\ \chi_{i_n} & \chi_{i_n$ |
|          | Thi Raz 0 0  |

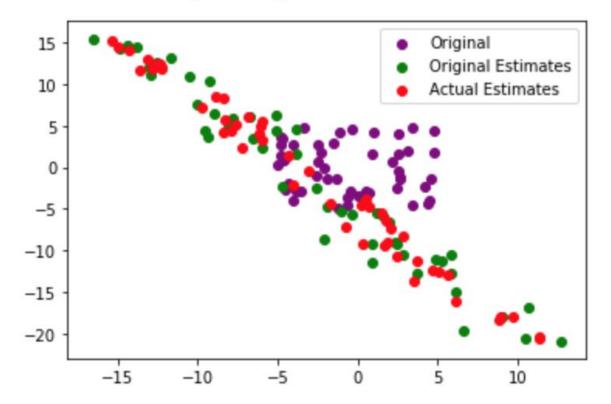
### 3) S & t matrices

#### T Matrix

```
array([[-1.87154926], [-3.05145812]])
```

#### **S Matrix**

4)
<matplotlib.legend.Legend at 0x13f3e7a90>

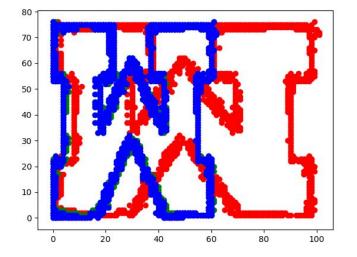


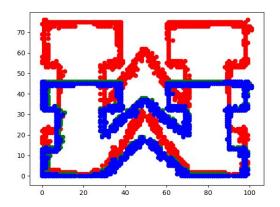
5) In order to transform the points, I take the dot product of the original X & Y points with my S\_matrix. For my new x value, I keep the first element from this dot product and for my new y value, I keep the second element from this dot product.

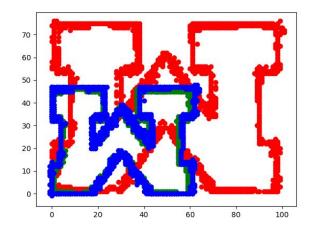
6)

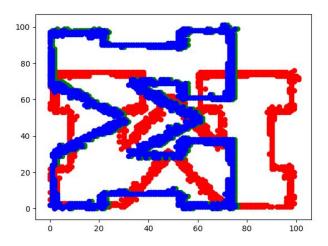
```
case 0
[[ 1.00555949e+00 1.61370672e-03 -1.35143989e-01]
  2.56045861e-03 6.22536404e-01 -7.35872070e-01]
 [ 4.51704286e-05 3.59823762e-05 1.00000000e+00]]
case
    - 1
[[ 6.20394776e-01 1.50050345e-03 -6.60977645e-01]
  4.62565656e-05 1.00232229e+00 -4.06369835e-02]
  3.66035139e-07
                  2.25931463e-05 1.00000000e+00]]
case 2
[[ 6.46216419e-01 1.01850154e-02 -1.45233663e+00]
  1.39156925e-02 6.46332905e-01 -1.60285342e+00]
                  2.76954685e-04 1.00000000e+00]]
 [ 3.11050183e-04
case 3
[[ 2.27055861e-14  1.00000000e+00
                                 2.85662774e-13]
  1.00000000e+00 -9.26614147e-16
                                 4.78080376e-14]
 [ 9.61481343e-17  0.00000000e+00
                                  1.00000000e+00]]
case 4
[[-1.00000000e+00 -1.45848347e-13
                                  7.60000000e+01]
  6.59624040e-13 -1.00000000e+00
                                 1.01000000e+021
 [ 9.52540533e-15  6.85280959e-16
                                 1.00000000e+00]]
case 5
[[ 4.01387046e-01 6.36777993e-01 -1.79445757e+00]
  6.40534201e-01 3.96745666e-01 -1.76059696e+00]
  2.21034153e-04 1.39932850e-04 1.00000000e+00]]
case 6
[[ 7.70056604e-01 4.54255201e-02 -6.34808816e-02]
 [-7.87253712e-02 8.63813194e-01 5.49671671e+00]
 [-3.05270237e-04 -6.37121535e-05 1.00000000e+00]]
case 7
[[ 8.41165566e-01 -4.86333708e-02
                                  4.30457199e+00]
  -4.77384685e-02 9.86999947e-01
                                  3.56055033e+00]
 [-4.55556648e-04 1.08171485e-04 1.00000000e+00]]
```

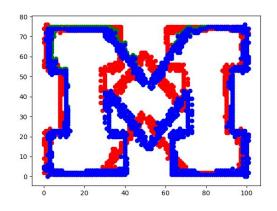
#### 7) Cases 0 through 8

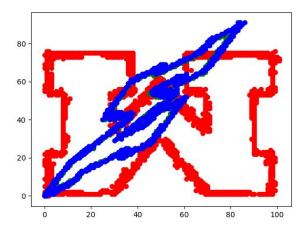


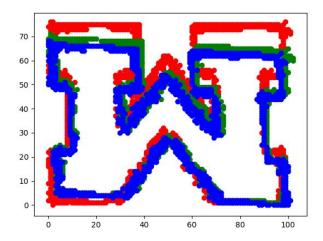












8) These transformations are unsurprising given the espoused function of a homography matrix. Obviously, this does not result in full coverage given that green is shown in many areas.

# #2 - Image Stitching

# 1) Grayscale images





## 2) Feature Points





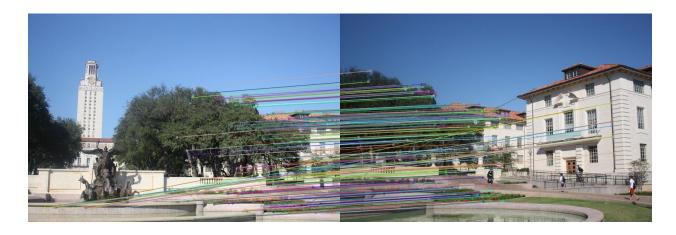
3) Distance was computed using Euclidean distance. Specifically:

```
distance = np.sqrt(np.sum((des1[:, np.newaxis, :] - des2[np.newaxis, :, :]) **
2, axis=-1))
```

4) The top 250 matches with the least distance were selected. Specifically, this function was used where the 'a' was the distance matrix and N was 250.

```
def smallestN_indices(a, N):
   idx = a.ravel().argsort()[:N]
   return np.stack(np.unravel_index(idx, a.shape)).T
```

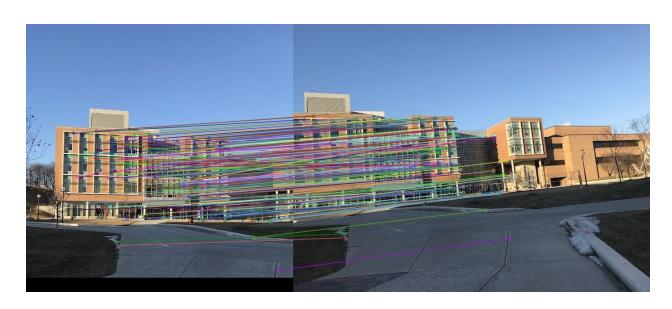
5) Matches  $\rightarrow$  199 inliers were found where the average distance of the residual was 97.0306. However a distance of less than 5 was required to qualify as an inlier.

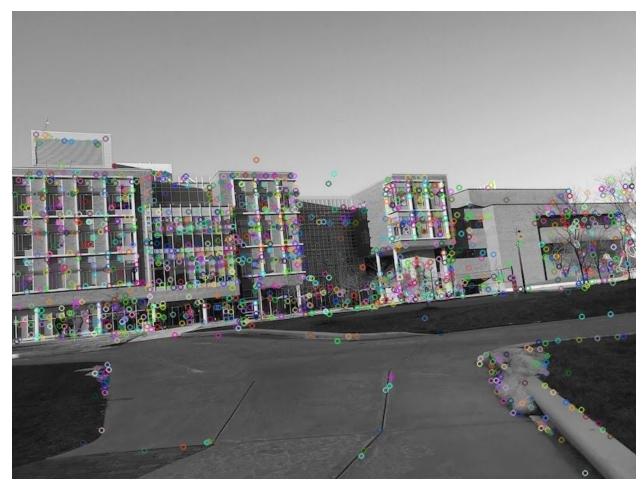


- 6)  $N/A \rightarrow see code$
- 7) Stitched image.



# 8) BBB image.







159 inliers with an average residual of 10.906617376795804.

