

# HW 3: HOMOGRAPHY AND FUNDAMENTAL MATRIX ESTIMATION

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## Output

And the output images are as follows...



Mapping between top 250 matched features



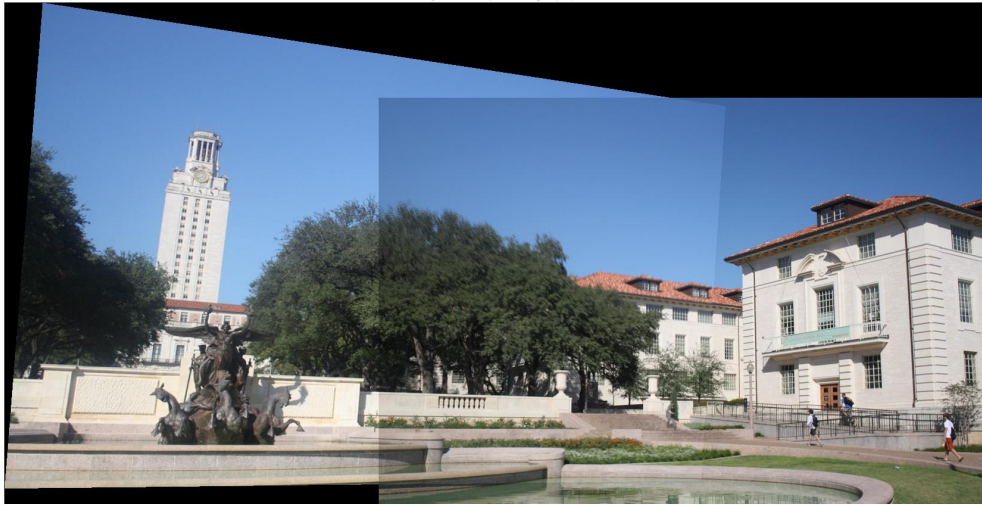
Transformed image



Number of inliers: 129

Average: 1.4125

Alignment by homography



The 129 inlier matches in both images



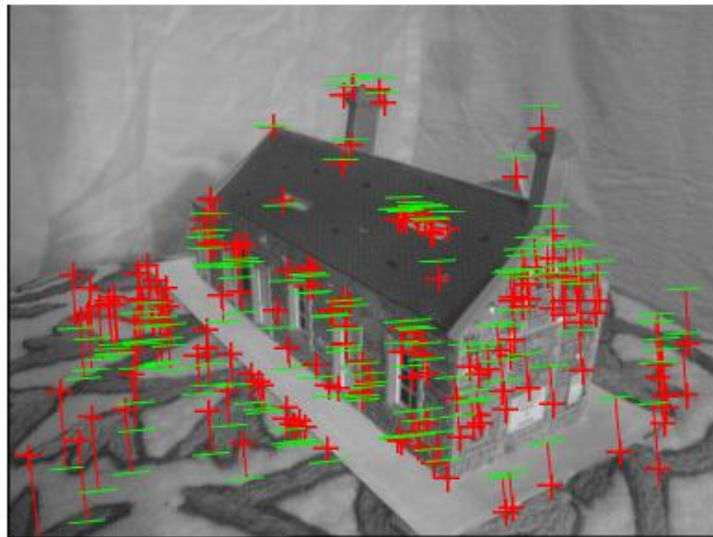
The matching between 129 inlier matches in both images

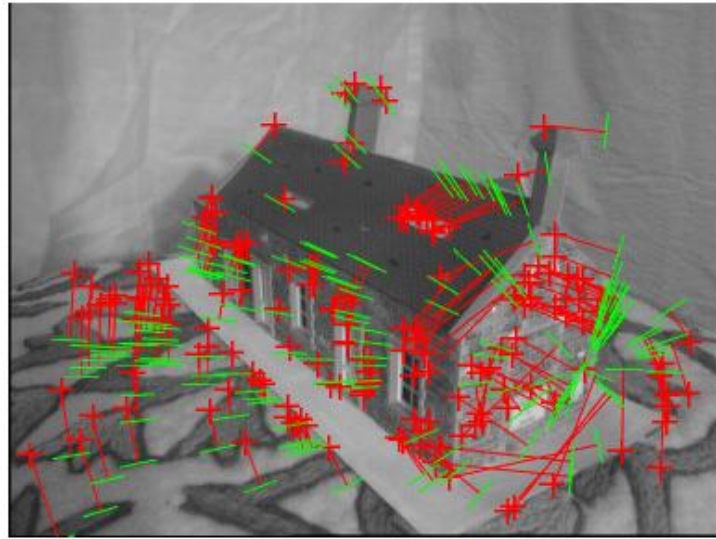






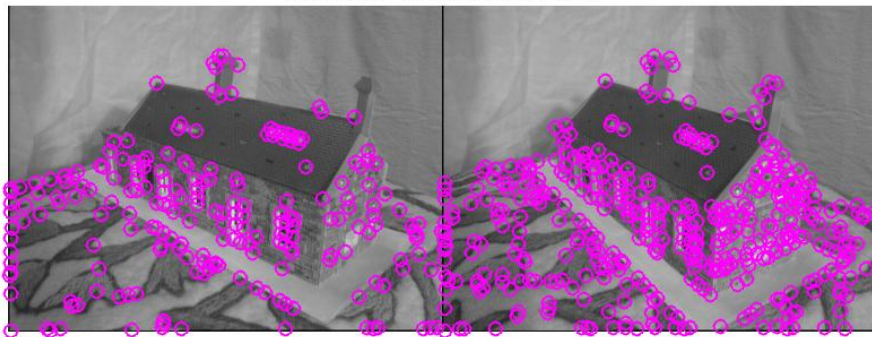
Normalised





Un-normalised

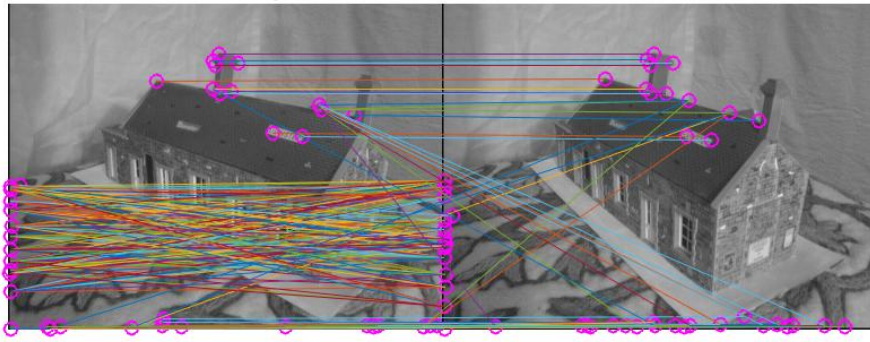
**Detected features/corners**



Top matched 300 features



Mapping between top 300 matched features



Number of inliers: 111

Average: 3.0563

