

# CVIP HW 3 REPORT

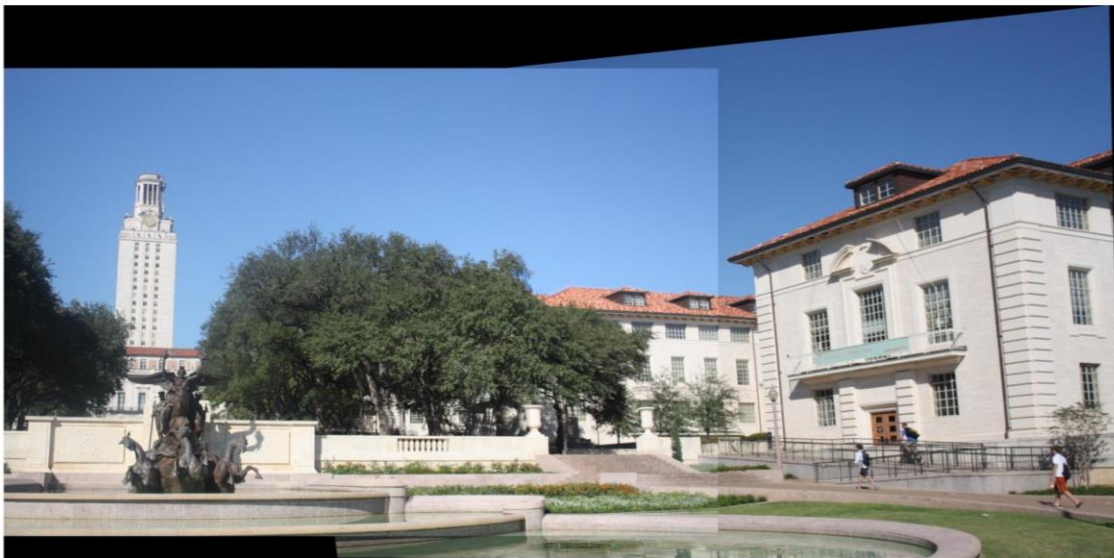
GAUTAM SHENDE 50245840

## 1) HOMOGRAPHY:

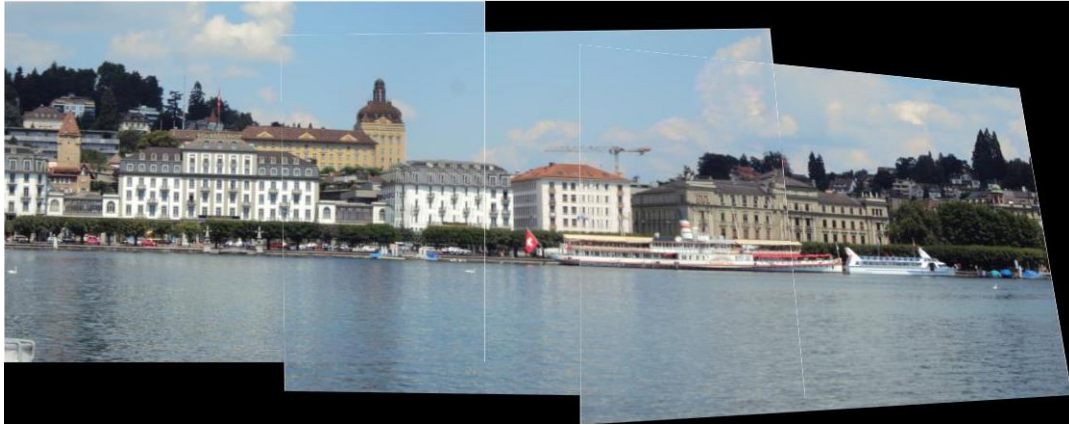
**A) Putative Matches:** I have used Blob detection from HW2 before getting SIFT descriptors)



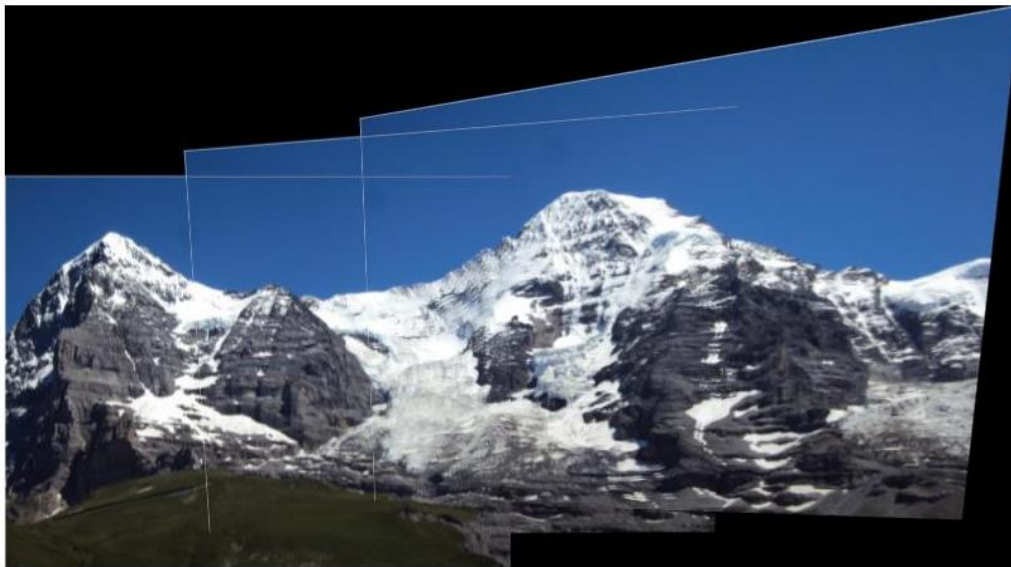
**B) Image Stitching Outputs including 3 image panoramas (Extra Credit)**



*Max\_inliers = 52, Mean inlier residual = 5.4384*



*Max\_inliers = 128, mean\_residual = 10.5469*



*Max\_inliers = 135, mean\_residual = 1.7555*



*Max\_inliers = 155, mean\_residual = 6.4514*

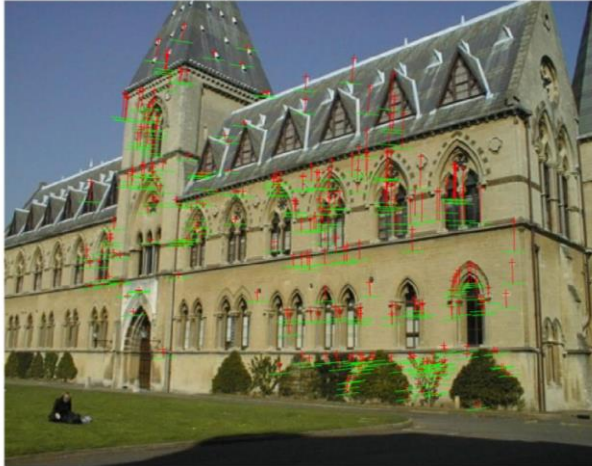


## 2.2) FUNDAMENTAL MATRIX & EPIPOLAR GEOMETRY

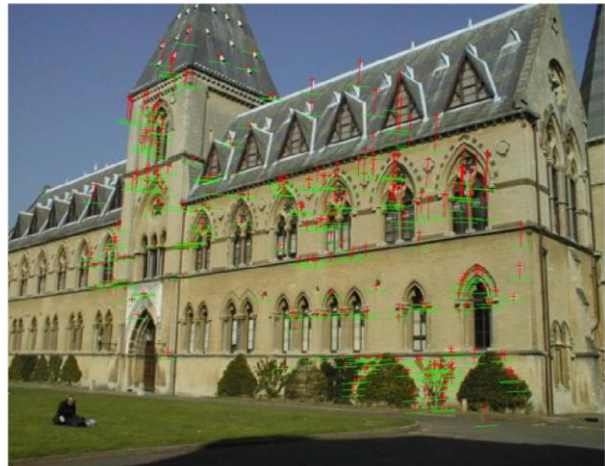
### GROUND TRUTH

Library Mean Residual Errors:

*Un-normalized: 11.8459*

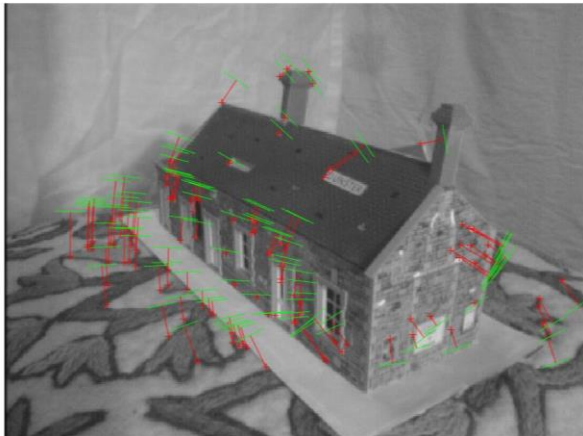


*Normalized: 9.9614*



House Mean Residual Errors:

*Un-normalized: 26.7532*



*Normalized: 14.1954*

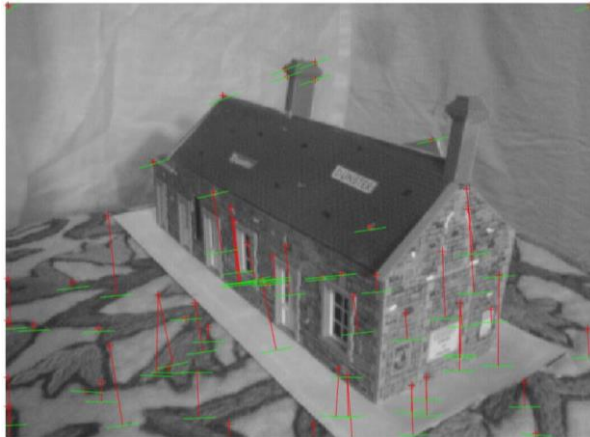


## 2.3) RANSAC

Since I used blobs in Part1, I decided to use Harris detection before getting SIFT descriptors in Part 2.

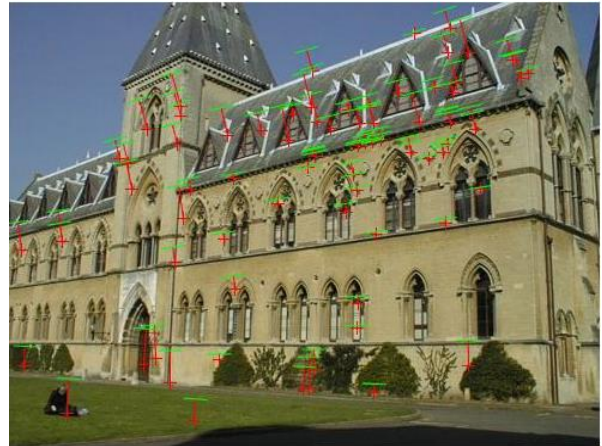
House Inlier Mean Residual error:

Threshold = 20, max inliers = 57/99, mean\_inliers\_res= 7.414



Library Inlier Mean Residual error:

Threshold = 30 max\_inliers = 93/199, mean\_inliers\_res =13.46



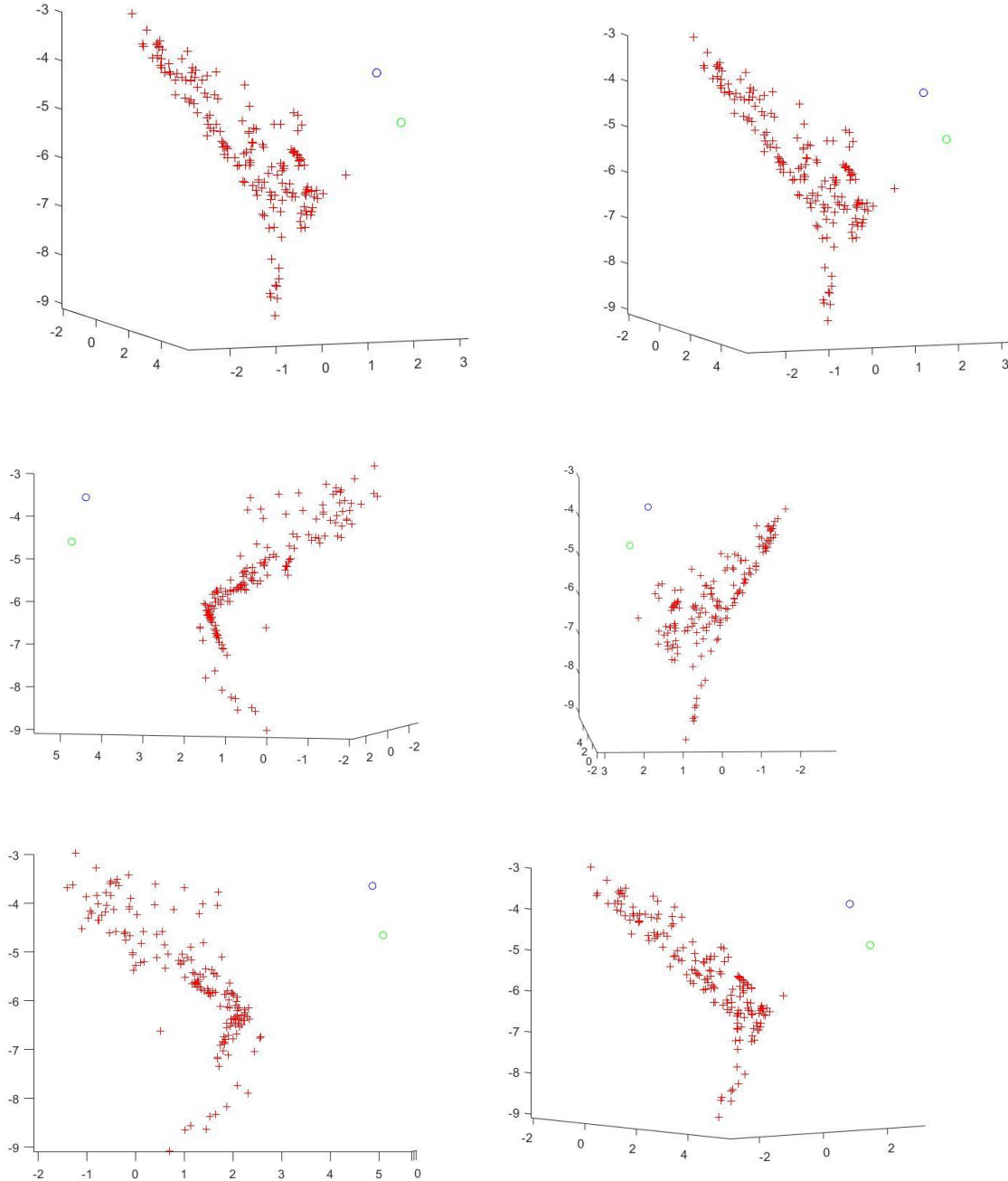
### ***Ground Truth results were much better in comparison to Ransac !***

With RANSAC, the general residual error had increased so I had to set a lower threshold in order to avoid erroneous results with overlapping epi-polar lines.

I was running RANSAC for a total of  $10^5$  iterations in the estimation of fundamental matrix and yet there was a lot of scope for improvement. The matches and epi-polar lines are far from accurate when compared to the ground truth matches. A lot of the parameters needed to be tuned in RANSAC based on the image. The library image needed more *correspondences* than the house.

## 2.4) Reconstructed Camera Centers and Points in 3D:

GIF : [https://drive.google.com/open?id=1CYs29KXmEc\\_Z8zJ0J0E2LOfGly2HbRpm](https://drive.google.com/open?id=1CYs29KXmEc_Z8zJ0J0E2LOfGly2HbRpm)



**Residual Errors between original 2D and projected 3D points:**

- Image1 = 0.0025,
- Image2 = 0.156

**References:**

*Homography and RANSAC:* [https://www.youtube.com/watch?v=oT9c\\_LIFBqs](https://www.youtube.com/watch?v=oT9c_LIFBqs)

*Fundamental Matrix:* <https://www.youtube.com/watch?v=DgGV3l82NTk>

*Image stitching:* <http://home.deib.polimi.it/boracchi/teaching/IAS/Stitching/stitch.html>

*Triangulation:* [dcyoung.weebly.com/fundamental-matrix--triangulation.html](http://dcyoung.weebly.com/fundamental-matrix--triangulation.html)