## **Lab 3 : computer vision**

# **Image Mosaics**

**April 27, 2019** 

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#### Overview

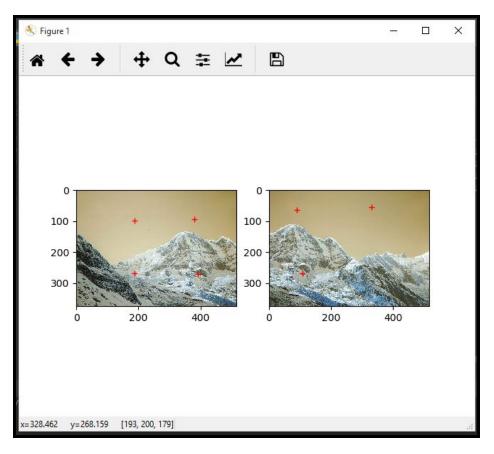
In this exercise, you will implement an image stitcher that uses image warping and homographies to automatically create an image mosaic. We will focus on the case where we have two input images that should form the mosaic, where we warp one image into the plane of the second image and display the combined views. This problem will give some practice manipulating homogeneous coordinates, computing homography matrices, and performing image warps. For simplicity, we will specify corresponding pairs of points manually using mouse clicks.

### Getting correspondence

#### Automatic Manual def auto\_correspondence(image1,image2): def manual\_correspondence(image1, image2, number\_of\_points): ' get coresspondance points between two given " Display images and select matching points " images using (oriented BRIEF) keypoint detector and fig = plt.figure() fig1 = fig.add\_subplot(1, 2, 1) descriptor extractor' ' better than SIFT descriptors' fig2 = fig.add subplot(1, 2, 2) orb = cv.ORB\_create() # Display the image keypoint1, descriptor1 = # to flip the image : #, origin='lower' orb.detectAndCompute(image1, None) fig1.imshow(image1) keypoint2, descriptor2 = fig2.imshow(image2) orb.detectAndCompute(image2, None) plt.axis('image') bf = cv.BFMatcher(cv.NORM HAMMING, p1 = np.zeros([(number of points // 2), 2]) crossCheck=True) # creates a matcher p2 = np.zeros([number\_of\_points // 2, 2]) pts = plt.ginput(n=number\_of\_points, # Match descriptors. timeout=0) matches = bf.match(descriptor1, descriptor2) # p1 itr = 0 matches the two descriptors p2\_itr = 0 matches = sorted(matches, key=lambda x: for i in range(0, number\_of\_points): x.distance) # sort matches where best matches come if i % 2 == 0: p1[p1\_itr] = pts[i] print("p1 of index ", p1\_itr, " is ", points1 = [] # list of correspondence points in pts[i]) p1\_itr += 1 first image points2 = [] # list of correspondence points in second image else: $p2[p2_itr] = pts[i]$ # loop on matches and fills points1 and points2 print("p2 of index ", p2 itr, " is ",

```
for match in matches:
    index1 = match.queryIdx
    points1
.append((int(keypoint1[index1].pt[0]),
int(keypoint1[index1].pt[1])))
    index2 = match.trainIdx
    points2.append((int(keypoint2[index2].pt[0]),
int(keypoint2[index2].pt[1])))

points1 = np.array(points1)
points2 = np.array(points2)
return points1, points2
```





### Computing the homography parameters

- 8 unknown variables
- we need at least 4 pair of points
- homography format is that the last element in the matrix is 1
- the resultant projective transformation is 3 X 3

```
A = np.zeros([2 * points1.shape[0], 8]) # construct A from input
points
    for i in range(points1.shape[0]): # constructs A from given points of
image 1
        A[i * 2] = points1[i, 0], points1[i, 1], 1, 0, 0, 0, -points1[i, 0]
* points2[i, 0], -points1[i, 1] * points2[i, 0]

        A[i * 2 + 1] = 0, 0, 0, points1[i, 0], points1[i, 1], 1, -points1[i, 0]
* points2[i, 1], -points1[i, 1] * points2[i, 1]

B = B.flatten().reshape(-1, 1) # flatten and reshape to be one column
for dimension suitability

H = np.linalg.lstsq(A, B, rcond=None)[0] # returns H

H = np.append(H, 1) # puts 1 at the end of H

H = np.reshape(H, [3, 3]) # return H as matrix of shape 3x3
return H
```

### Warping between image planes

- forward warping
- The transformed coordinates are sub-pixel values, sample the pixel values from nearby pixels.
- For color images, warp each RGB channel separately and then stack together to form the output
- inverse warping to remove holes

## Screenshots of the output mosaic

Manual : result\_manual with ransac\_ransac\_loops=10













Warp one image into a frame region in the second image. To do this, let the points
from the one view be the corners of the image you want to insert in the frame, and let the
corresponding points in the second view be the clicked points of the frame
(quadrilateral) into which the first image should be warped





#### Extra credits section

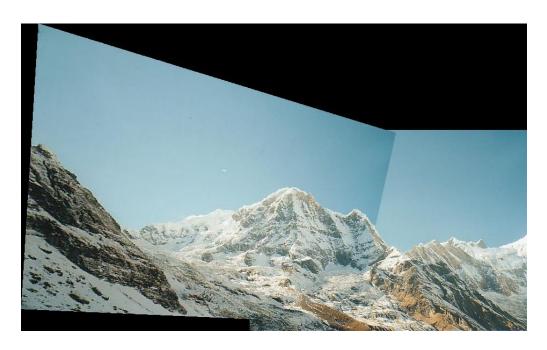
#### RANSAC

```
def ransac(points1, points2, threshold, iterations):
    "takes correspondence points and for each 4 random pairs it
calculates h "
    " and counts inliners from a given threshold and keeps the
best h"
    max inliners = 0
    best_h = None
    for i in range(iterations):
        inliners = 0
        randp = np.zeros([4, 2])
        randp_ = np.zeros([4, 2])
        H = None
        for j in range(4):
            random_index = random.randrange(0,
points1.shape[0], 1) # picks random points from the given set
            randp[j] = points1[random_index]
            randp_[j] = points2[random_index]
        H = Homography.calculate_homography(randp, randp_) #
calculates h from the 4 random points
        for j in range(points1.shape[0]):
            error = ransac_error(points1[j], points2[j], H)
            if (error < threshold):</pre>
                inliners += 1
        if (inliners > max_inliners):
            max_inliners = inliners
            best h = H
            # return the H and the number of inliners for H
    return best_h, max_inliners
```

• MANUAL\_CORRESPONDENCE, mosaic\_builder.RANSAC\_H,10

shown before

• AUTO\_CORRESPONDENCE, mosaic\_builder.RANSAC\_H, 100





### AUTO\_CORRESPONDENCE, mosaic\_builder.RANSAC\_H, 1000



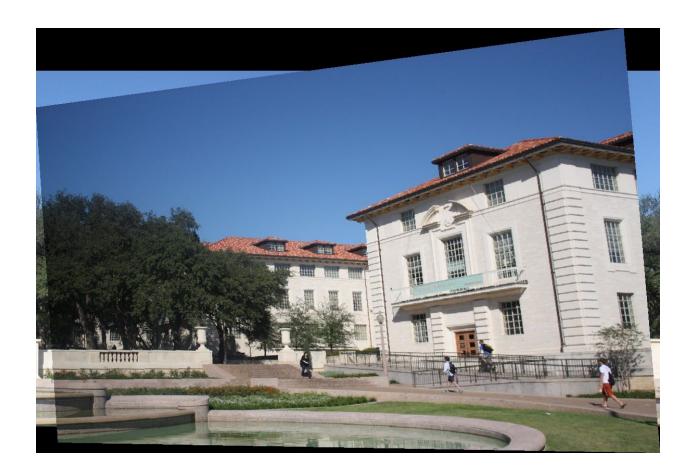


### AUTO\_CORRESPONDENCE, mosaic\_builder.RANSAC\_H, 5000





# AUTO\_CORRESPONDENCE, mosaic\_builder.NORMAL\_H (without RANSAC)



### Automatic correspondence shown before

#### • Conclusion:

the first pair of images couldn't be formed under automatic correspondence without RANSAC

it robustly estimates the homography matrix from noisy correspondences and it successfully gives good results even when there are outlier (bad) correspondences given as input