HW2 Image stitching

1. Explain your implementation

In this homework, I follow the architecture provided by TA.

• Detecting key point(feature) on the images

I use the SIFT algorithm to find keypoints and descriptors of images. The method I use to implement is to call function provided by OpenCV. Since SIFT algorithm only can handle gray image, I preprocess them into grayscale before using it.

```
img1,img_gray1 = read_img(img_path_lists[0])
img2,img_gray2 = read_img(img_path_lists[1])
kts1, dcpt1 = SIFT_detector.detectAndCompute(img_gray1, None)
kts2, dcpt2 = SIFT_detector.detectAndCompute(img_gray2, None)
```

Finding features correspondences (feature matching)

To find the corresponding key points in a paired image, we utilize the KNN algorithm. We implement it using brute force. Besides, we use Lowe's Ratio test algorithm to filter out some bad matching.(Threshold = 0.75)

```
def feature_matching_implement(kts1, dcpt1,kts2, dcpt2):
    good_matches = []
    for ii,des1 in enumerate(tqdm(dcpt1)):
    min_value_1_distance = 100000000000000001
         min_value_2_idx = -1
for jj,des2 in enumerate(dcpt2):
             distance = (des1 - des2) * (des1 - des2)
distance = sum(distance)
             distance = math.sqrt(distance)
             if(distance < min value 1 distance):</pre>
                  min_value_2_distance = min_value_1_distance
                  min_value_2_idx = min_value_1_idx
                  min value 1 distance = distance
                  min value 1 idx = jj
             elif(distance < min_value_2_distance and distance >= min_value_1_distance):
    min_value_2_distance = distance
         min_value_2_idx = jj
if(min_value_1_distance < min_value_2_distance*0.75):</pre>
             m = cv2.DMatch()
             m.queryIdx = ii
             m.trainIdx = min_value_1_idx
m.distance = min_value_1_distance
             m.imgIdx = 0
              good_matches.append(m)
    return good_matches
```

Computing homography matrix.

Because of the existence of outliers, we utilize the RANSAC algorithm to find an appropriate homography that can transform pixels on image A to imageB. In the RANSAC algorithm, we randomly select four paired points and use them to estimate the homography matrix.

After getting an estimated homography matrix, we apply this matrix to all key points to get the size of the consensus set. If this consensus set is larger than ever, we mark this homography matrix as "best homography matrix". We repeat this process MAX_ITERATION times.

Stitching image (warp images into same coordinate system)

In this section we wrap images into the same coordinate system. We can use "best homography matrix" to transform pixels from image A to image B. However, some pixels from image A may translate to values outside the boundary of image B, either less than 0 or exceeding its own boundary. Thus, we construct another matrix A to ensure that all translated pixel positions will be greater than 0 and determine the size of transformed image by the translated corners of image A.

There still remain two questions to solve, one for blendering and the other for gain compensations.

❖ Blendering:

Since our homography matrix is not perfect, it is weird if we put two images together directly.



We could use some blendering algorithm to help. The algorithm I use is "linearBlendingWithConstantWidth". In my implementation, I use the following equation:

$$Image_{result} = \alpha * image_A + (1 - \alpha) * image_B$$

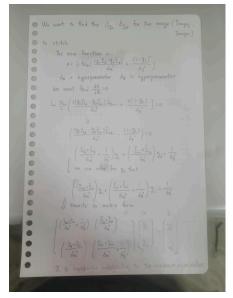
This equation is only applied to a constant width in the overlap area.

```
def HamorelandingwithConstantWidth(solf, logs):

| linear Elending with Constant Width, worlding ghost region
| a you need to determine the size of constant with
| say_inft, ang_right = sizes
| say_inft, say_inft, say_inft
| say_inft
|
```

Gain compensations:

The Gain compensations want to solve the problem for the difference of Intensity between two different paired images. I derive the equation that helps me to construct a matrix that optimizes the result.



Follow the the result of derivation, we can find the best $\sigma_{\!{}_A}$ and $\sigma_{\!{}_B}.$

```
image * np.repeat(overlap[:, :, np.nesexts], 3, exts=2),
exts=(0, 1),
t_gain_compensations(
rpl,warpl,sigma_n: float = 10.0, sigma_g: float = 0.1
 fs = { np.zeros(1) for _ in range(2) }
ult = np.zeros(1)
fs(0) = {
(1 / zigns_n ** 2) * pair_natch_I_0 * pair_natch_I_1.
```

- 2. Show the result of stitching "Base" images (and "Challenge image" if you did that part).
 - Base images



• Challenge images



3. Discuss different blending method results.

I have tried the following method to blend images.

Linear blending vs Linear blending with constant width
 The result of these two methods is not significant. However, Linear blending
 with constant width can save lots of computation resources compared to
 Linear blending. Because Linear blending with constant width only constructs
 the alpha part in constant width instead constructing in the whole overlap
 region.



The left is the alpha map in "Linear blending with constant width". The middle part is the alpha map in "Linear blending". The right part is the overlap region of two images.

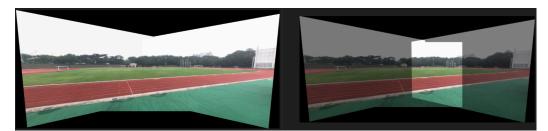


The result of Linear blending and Linear blending with constant width. The left one is Linear blending. The right one is Linear blending with constant width.

Simple blending vs Multiband blending
 Compared to simple blending, multi band blending needs more computational
 time and resources. Besides, the result of multi band blending is more blur.
 However, the result of multi band blending is more seemless.



This is the result of simple blending. The left is the whole result. The right part focuses on the overlap region.



This is the result of multiband blending. The left is the whole result. The right part focuses on the overlap region.