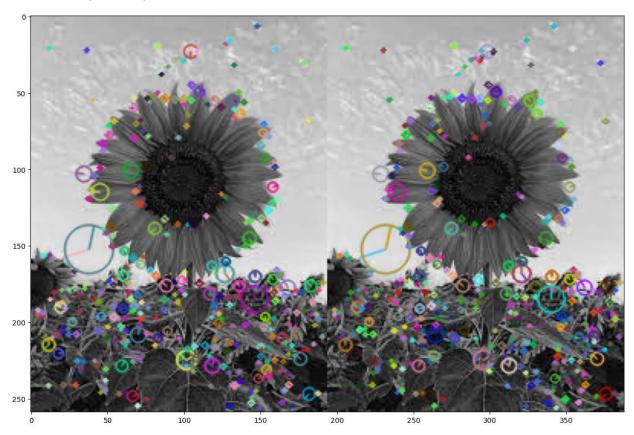
```
In [1]: import cv2
        import matplotlib.pyplot as plt
        import numpy as np
        import random
        from tqdm.notebook import tqdm
        plt.rcParams['figure.figsize'] = [15, 15]
In [2]: # Read image and convert them to gray!!
        def read_image(img):
            img = cv2.imread('F:\VIT BHOAPL\Fall Sem 23-24\Deep Learning\deep learning activities\RANSAC ALgo\sunflower2.jpg
            img_gray= cv2.cvtColor(img, cv2.COLOR_RGB2GRAY)
            img_rgb = cv2.cvtColor(img, cv2.COLOR_BGR2RGB)
            return img_gray, img, img_rgb
In [3]: left_gray, left_origin, left_rgb = read_image('data/1.jpg')
        right_gray, right_origin, right_rgb = read_image('data/2.jpg')
In [4]: def SIFT(img):
            siftDetector= cv2.xfeatures2d.SIFT_create() # limit 1000 points
            # siftDetector= cv2.SIFT_create() # depends on OpenCV version
            kp, des = siftDetector.detectAndCompute(img, None)
            return kp, des
        def plot_sift(gray, rgb, kp):
            tmp = rgb.copy()
            img = cv2.drawKeypoints(gray, kp, tmp, flags=cv2.DRAW_MATCHES_FLAGS_DRAW_RICH_KEYPOINTS)
            return img
In [5]: # Better result when using gray
        kp_left, des_left = SIFT(left_gray)
        kp_right, des_right = SIFT(right_gray)
In [6]: kp_left_img = plot_sift(left_gray, left_rgb, kp_left)
        kp_right_img = plot_sift(right_gray, right_rgb, kp_right)
        total_kp = np.concatenate((kp_left_img, kp_right_img), axis=1)
```

Out[6]: <matplotlib.image.AxesImage at 0x7906eaf7ebc0>

plt.imshow(total_kp)



```
In [7]:
def matcher(kp1, des1, img1, kp2, des2, img2, threshold):
    # BFMatcher with default params
    bf = cv2.BFMatcher()
    matches = bf.knnMatch(des1,des2, k=2)

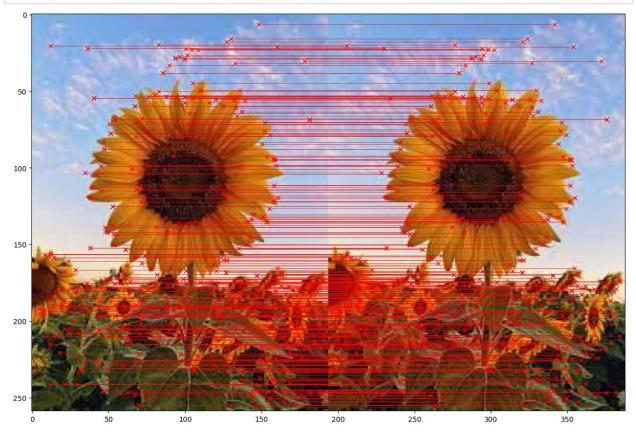
# Apply ratio test
good = []
for m,n in matches:
    if m.distance < threshold*n.distance:
        good.append([m])

matches = []
for pair in good:
    matches.append(list(kp1[pair[0].queryIdx].pt + kp2[pair[0].trainIdx].pt))

matches = np.array(matches)
return matches</pre>
```

```
In [8]: matches = matcher(kp_left, des_left, left_rgb, kp_right, des_right, right_rgb, 0.5)
```

In [10]: total_img = np.concatenate((left_rgb, right_rgb), axis=1) plot_matches(matches, total_img) # Good mathces



```
In [11]: def homography(pairs):
              rows = []
              for i in range(pairs.shape[0]):
                  p1 = np.append(pairs[i][0:2], 1)
                  p2 = np.append(pairs[i][2:4], 1)
                  row1 = [0, 0, 0, p1[0], p1[1], p1[2], -p2[1]*p1[0], -p2[1]*p1[1], -p2[1]*p1[2]]
row2 = [p1[0], p1[1], p1[2], 0, 0, 0, -p2[0]*p1[0], -p2[0]*p1[1], -p2[0]*p1[2]]
                  rows.append(row1)
                  rows.append(row2)
              rows = np.array(rows)
              U, s, V = np.linalg.svd(rows)
              H = V[-1].reshape(3, 3)
              H = H/H[2, 2] # standardize to let w*H[2,2] = 1
              return H
In [12]: def random_point(matches, k=4):
              idx = random.sample(range(len(matches)), k)
              point = [matches[i] for i in idx ]
              return np.array(point)
In [13]: def get_error(points, H):
              num_points = len(points)
              all_p1 = np.concatenate((points[:, 0:2], np.ones((num_points, 1))), axis=1)
              all_p2 = points[:, 2:4]
              estimate_p2 = np.zeros((num_points, 2))
              for i in range(num_points):
                  temp = np.dot(H, all_p1[i])
                  estimate_p2[i] = (temp/temp[2])[0:2] # set index 2 to 1 and slice the index 0, 1
              errors = np.linalg.norm(all_p2 - estimate_p2 , axis=1) ** 2
              return errors
In [14]: def ransac(matches, threshold, iters):
              num_best_inliers = 0
              for i in range(iters):
                  points = random_point(matches)
                  H = homography(points)
                  # avoid dividing by zero
                  if np.linalg.matrix_rank(H) < 3:</pre>
                       continue
                  errors = get_error(matches, H)
                  idx = np.where(errors < threshold)[0]</pre>
                  inliers = matches[idx]
                  num_inliers = len(inliers)
                  if num_inliers > num_best_inliers:
                      best_inliers = inliers.copy()
                      num_best_inliers = num_inliers
                      best_H = H.copy()
              print("inliers/matches: {}/{}".format(num_best_inliers, len(matches)))
              return best_inliers, best_H
```

In [15]: inliers, H = ransac(matches, 0.5, 2000)

inliers/matches: 435/435