## Assignment 2 - EN3160

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Github Link https://github.com/RPX2001/Assignment2\_EN3160.git

#### Question 1 - Blob Detection

```
def generate_log_kernel(sigma):
       """Generate Laplacian of Gaussian (LoG) kernel."""
      radius = int(3 * sigma)
      x, y = np.meshgrid(np.arange(-radius, radius + 1), np.arange(radius, radius + 1))
      kernel = ((x**2 + y**2) / (2 * sigma**2) - 1) * np.exp(-(x**2 + y**2) / (2 *
          sigma**2)) / (np.pi * sigma**4)
      return kernel
  def find_blob_centers(log_image, sigma):
      """Find local maxima in the LoG response image."""
      coords = []
      height, width = log_image.shape
      neighborhood = 1  # 3x3 window size
      for row in range(neighborhood, height-neighborhood):
          for col in range(neighborhood, width-neighborhood):
               local_window = log_image[row-neighborhood:row+neighborhood+1, col-
                  neighborhood:col+neighborhood+1]
               max_response = np.max(local_window)
               if max_response >= 0.10: # Adjust threshold for detection
                   x_offset , y_offset = np.unravel_index(np.argmax(local_window),
                      local_window.shape)
                   coords.append((row + x_offset - neighborhood, col + y_offset -
18
                      neighborhood))
      return set(coords)
```

To optimally detect a blob with radius r, the relationship was used.sigma values corresponding to r in the range of (1, 10) were applied for blob detection. This approach ensures that the scale parameter sigma is appropriately tuned for detecting blobs of different sizes, corresponding to the radi in the given range.





Figure 1: Blob Detection

#### Question 2 - Line and circle fitting using RANSAC

```
def line_distance(points, a, b, d):
   return np.abs(a * points[:, 0] + b * points[:, 1] + d) / np.sqrt(a**2 + b**2)
   def fit_line(p1, p2):
       a = p1[1] - p2[1]
                          # y1 - y2
       b = p2[0] - p1[0] # x2 - x1
       d = -(a * p1[0] + b * p1[1])
                                      \# ax1 + by1 + d = 0
       norm = np.sqrt(a**2 + b**2)
       return a / norm, b / norm, d / norm
   # RANSAC for line fitting
   def ransac_line(points, threshold, num_iterations=100):
10
       best_a, best_b, best_d = None, None, None
       best_consensus = []
       for _ in range(num_iterations):
           # Randomly select 2 points
14
           p1, p2 = points[sample(range(len(points)), 2)]
           a, b, d = fit_line(p1, p2)
16
           distances = line_distance(points, a, b, d)
           consensus_set = points[distances < threshold]</pre>
           if len(consensus_set) > len(best_consensus):
19
               best_a, best_b, best_d = a, b, d
               best_consensus = consensus_set
       return best_a, best_b, best_d, best_consensus
   def circle_distance(points, x0, y0, r):
       return np.abs(np.sqrt((points[:, 0] - x0)**2 + (points[:, 1] - y0)**2) - r)
24
   def fit_circle(p1, p2, p3):
       A = np.array([[p1[0], p1[1], 1],
26
                      [p2[0], p2[1], 1],
                      [p3[0], p3[1], 1]])
28
       B = np.array([-(p1[0]**2 + p1[1]**2),
29
                      -(p2[0]**2 + p2[1]**2),
30
                      -(p3[0]**2 + p3[1]**2)])
31
32
       sol = np.linalg.solve(A, B)
       x0, y0 = -0.5 * sol[0], -0.5 * sol[1]
33
       r = np.sqrt((sol[0]**2 + sol[1]**2) / 4 - sol[2])
34
       return x0, y0, r
35
   def ransac_circle(points, threshold, num_iterations=100):
36
       best_x0, best_y0, best_r = None, None, None
37
       best_consensus = []
38
       for _ in range(num_iterations):
39
           # Randomly select 3 points
40
           p1, p2, p3 = points[sample(range(len(points)), 3)]
41
           x0, y0, r = fit_circle(p1, p2, p3)
42
           distances = circle_distance(points, x0, y0, r)
43
           consensus_set = points[distances < threshold]</pre>
44
           if len(consensus_set) > len(best_consensus):
45
               best_x0, best_y0, best_r = x0, y0, r
46
               best_consensus = consensus_set
47
       return best_x0, best_y0, best_r, best_consensus
48
   threshold_line = 0.5
   threshold_circle = 0.5
  a, b, d, line_consensus = ransac_line(X, threshold_line)
  X_remnant = np.array([point for point in X if point not in line_consensus])
  x0, y0, r, circle_consensus = ransac_circle(X_remnant, threshold_circle)
```

This approach ensures that the scale parameter sigma is appropriately tuned for detecting blobs of different sizes, corresponding to the radii in the given range.

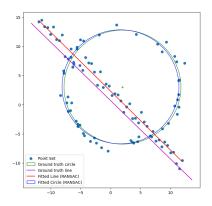


Figure 2: Line and Circle fitting









Figure 3: Superimposed image

### **Question 3 - Superimposing an image on another**

```
def superimpose(image, logo, dst_points, beta=0.3, alpha=1):
    h, w, _ = logo.shape
    src_points = np.array([(0, 0), (w, 0), (w, h), (0, h)], dtype=np.float32)
    tform = transform.estimate_transform('projective', src_points, dst_points)
    tf_img = transform.warp(logo, tform.inverse, output_shape=image.shape[:2])
    tf_img = (tf_img * 255).astype(np.uint8)
    mask = np.any(tf_img > 0, axis=-1).astype(np.uint8)
    blended = cv.addWeighted(image, alpha, tf_img, beta, 0)
    result = np.where(mask[:, :, np.newaxis], blended, image)
    return result
```

# Question 4 – Image stitching

There were insufficient matching features between images 1 and 5 to accurately compute a homography directly. As a result, homographies were calculated between consecutive image pairs, such as 1-2, 2-3, and so on, where the images had more similarities. Image 1 was then progressively transformed through each of these computed homographies to align it with image 5.

```
def extract_SIFT_features(image1, image5, display = False):
    gray_img1 = cv.cvtColor(image1, cv.COLOR_RGB2GRAY)
    gray_img5 = cv.cvtColor(image5, cv.COLOR_RGB2GRAY)
    sift_detector = cv.SIFT_create(nOctaveLayers=3, contrastThreshold=0.09,
        edgeThreshold=25, sigma=1)
    kp1, desc1 = sift_detector.detectAndCompute(gray_img1, None)
    kp5, desc5 = sift_detector.detectAndCompute(gray_img5, None)
    matcher = cv.BFMatcher()
```









Figure 4: SIFT Matching

```
knn_matches = matcher.knnMatch(desc1, desc5, k=2)
8
       valid_matches = []
       for m, n in knn_matches:
           if m.distance < 0.75 * n.distance:</pre>
                valid_matches.append(m)
       return valid_matches, kp1, kp5
14
15
   def identify_inliers(source_pts, destination_pts, homography, threshold):
16
       projected_pts = homography(source_pts)
       distances = np.sqrt(np.sum((projected_pts - destination_pts) ** 2, axis=1))
18
19
       return np.where(distances < threshold)[0]</pre>
20
   def compute_best_homography(valid_matches, kp1, kp5):
       src_points_set = []
       dst_points_set = []
24
       for match in valid_matches:
25
           src_points_set.append(np.array(kp1[match.queryIdx].pt))
26
           dst_points_set.append(np.array(kp5[match.trainIdx].pt))
28
       src_points_set = np.array(src_points_set)
29
       dst_points_set = np.array(dst_points_set)
30
       ransac_points = 4
31
       error\_threshold = 1
32
       min_inliers = 0.5 * len(valid_matches)
33
       max_iterations = 200
34
       best_homography = None
35
       max_inliers = 0
36
       best_inliers_indices = None
37
       for i in range(max_iterations):
38
           random_matches = np.random.choice(valid_matches, ransac_points, replace=False
39
               )
           src_ransac = []
40
           dst_ransac = []
41
           for match in random_matches:
42
                src_ransac.append(np.array(kp1[match.queryIdx].pt))
43
                dst_ransac.append(np.array(kp5[match.trainIdx].pt))
44
           src_ransac = np.array(src_ransac)
45
           dst_ransac = np.array(dst_ransac)
46
           homography_estimation = transform.estimate_transform('projective', src_ransac
47
               , dst_ransac)
           inliers_indices = identify_inliers(src_points_set, dst_points_set,
48
               \verb|homography_estimation|, error_threshold|)
           if len(inliers_indices) > max_inliers:
49
                max_inliers = len(inliers_indices)
50
                best_homography = homography_estimation
51
                best_inliers_indices = inliers_indices
52
       print(f'Maximum_inliers_count_=[max_inliers]')
53
       return best_homography, best_inliers_indices
```