

# EN3160 Assignment 2- Fitting

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GitHub : <https://github.com/DinethraDivanjana2001/EN3166-Image-processing-and-machine-vision/tree/main/A02%20-%20Fitting>

## 1.0. Question 01-Blob Detection in a Sunflower Field Image

First, I converted the sunflower field image to grayscale to simplify blob detection. The algorithm applies Gaussian blur followed by a Laplacian filter to detect circular shapes at multiple scales.

- **Gaussian Blur:** Smooths the image, reducing noise.
- **Laplacian Filter:** Identifies circular edges by detecting intensity changes.

I varied the scale values ( $\sigma$ ) between 1.0 and 4.0 in 8 equal steps, as larger scales can capture larger structures, while smaller scales highlight finer details.

The range of  $\sigma$  values used was from 1.0 to 4.0

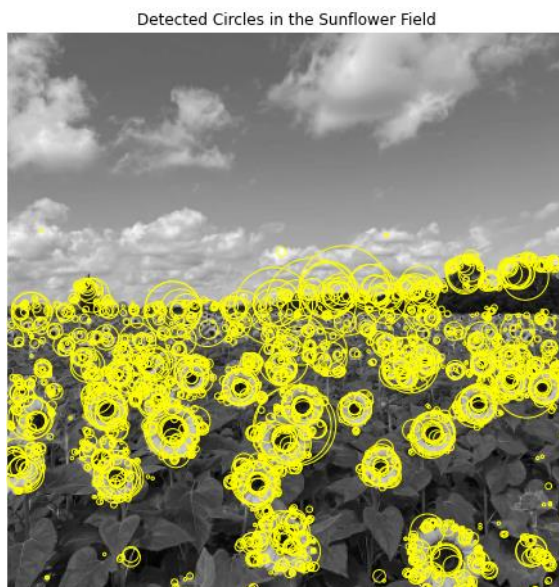


Figure 1-Detected Circles

```
Largest Blob Parameters:  
Center: (216, 166)  
Radius: 29  
Scale: 1.9192898346492004
```

Figure 2 - parameters of the largest circle

```
def apply_gaussian_laplacian(image, sigma_value):  
    """Apply Gaussian Blur followed by Laplacian filtering."""  
    blurred = cv.GaussianBlur(image, (0, 0), sigma_value)  
    laplacian = cv.Laplacian(blurred, cv.CV_64F)  
    return np.abs(laplacian)
```

Figure 2 - Gaussian-Laplacian Transformation

```
def detect_blobs(image, min_scale, max_scale, num_scales, threshold):  
    """Detect blobs across multiple scales and return the detected blobs."""  
    scales = np.linspace(min_scale, max_scale, num_scales)  
    blobs = []  
  
    for scale in scales:  
        adjusted_scale = scale / np.sqrt(2)  
        print(f"Processing scale: {adjusted_scale}")  
  
        laplacian_image = apply_gaussian_laplacian(image, adjusted_scale)  
  
        binary_mask = laplacian_image > threshold * laplacian_image.max()  
  
        contours, _ = cv.findContours(binary_mask.astype(np.uint8),  
                                     cv.RETR_EXTERNAL,  
                                     cv.CHAIN_APPROX_SIMPLE)  
  
        for contour in contours:  
            if len(contour) >= 5:  
                (x, y), radius = cv.minEnclosingCircle(contour)  
                blobs.append((int(x), int(y), int(radius), adjusted_scale))  
  
    return sorted(blobs, key=lambda b: -b[1])
```

Figure 4 - detects blobs by applying the Gaussian-Laplacian transformation across scales and extracting circles from contours

## 2.0. Question - 02

Implemented the RANSAC algorithm to estimate a line and a circle from a noisy set of points.

For line estimation, I constrained the line parameters  $[a, b]$  to  $[a, b]$ . I selected the error as the perpendicular distance from points to the line and defined a threshold for consensus, ensuring only points within this distance were considered in the fitting process. After obtaining the best-fit line, I subtracted the consensus points to analyse the remaining points for circle fitting. Using RANSAC again, I calculated the radial error for circle fitting and set a threshold to determine consensus points for the circle. Finally, I visualized the results by plotting the point set, the estimated line and circle, their inliers, and the sample points used for their estimation, presenting a comprehensive view of the fitting results.

- The `calculate_line_equation` function computes normalized line coefficients from two points; `total_least_squares_error` calculates the sum of squared errors for a line model; `constraint_function` enforces the normalization constraint  $a^2 + b^2 = 1$ ; and `identify_inliers` determines inliers by comparing the absolute error to a threshold. (figure 5)
- The `compute_circle_from_points` function calculates the center and radius of a circle given three points; `identify_circle_inliers` identifies inliers based on the radial error being below one-third of the circle's radius; `get_random_sample` selects three random points from a list for circle fitting; `calculate_distance_from_center` computes the distance of points from the circle's center; `algebraic_distance` determines the deviation of point distances from the mean radius; and `fit_circle_via_least_squares` fits a circle using least-squares optimization to estimate the center and radius. (figure 6)

```
Ratio of Inliers = 7.017543859649122 %
Inliers array has insufficient dimensions for plotting.
Center of RANSAC Circle = (2.0435741271449093, 3.870549248953072)
Radius of RANSAC Circle = 9.432686503699676
```

```
def calculate_line_equation(point1_x, point1_y, point2_x, point2_y):
    """Calculate line equation coefficients from two points."""
    delta_x = point2_x - point1_x
    delta_y = point2_y - point1_y
    magnitude = math.sqrt(delta_x**2 + delta_y**2)
    coeff_a = delta_y / magnitude
    coeff_b = -delta_x / magnitude
    intercept = (coeff_a * point1_x) + (coeff_b * point1_y)
    return coeff_a, coeff_b, intercept

def total_least_squares_error(coefficients, selected_indices):
    """Calculate total least squares error for a line model."""
    coeff_a, coeff_b, intercept = coefficients
    return np.sum(np.square(coeff_a * data_points[selected_indices, 0] +
                             coeff_b * data_points[selected_indices, 1] - intercept))

def constraint_function(coefficients):
    """Define constraint for coefficients."""
    return coefficients[0]**2 + coefficients[1]**2 - 1

constraints = ({'type': 'eq', 'fun': constraint_function})

def identify_inliers(data, coefficients, threshold):
    """Identify inliers based on the consensus function."""
    coeff_a, coeff_b, intercept = coefficients
    error = np.absolute(coeff_a * data[:, 0] + coeff_b * data[:, 1] - intercept)
    return error < threshold
```

Figure 5

```
def compute_circle_from_points(points):
    """Calculate the center and radius of the circle from three points"""
    point1, point2, point3 = points[0], points[1], points[2]
    temp = point2[0]**2 + point2[1]**2
    half_bc = (point1[0]**2 + point1[1]**2 - temp) / 2
    half_cd = (temp - point3[0]**2 - point3[1]**2) / 2
    determinant = (point1[0] - point2[0]) * (point2[1] - point3[1]) - (point2[0] - point3[0]) * (point1[1] - point2[1])

    # Center of the circle
    center_x = (half_bc * (point2[1] - point3[1]) - half_cd * (point1[1] - point2[1])) / determinant
    center_y = ((point1[0] - point2[0]) * half_cd - (point2[0] - point3[0]) * half_bc) / determinant

    radius = np.sqrt((center_x - point1[0])**2 + (center_y - point1[1])**2)
    return ((center_x, center_y), radius)

def identify_circle_inliers(points_list, center, radius):
    """Identify inliers to a circle model based on a threshold"""
    inliers = []
    threshold = radius / 3 # Set threshold to one-third of the radius

    for point in points_list:
        distance_error = np.sqrt((point[0] - center[0])**2 + (point[1] - center[1])**2) - radius
        if distance_error < threshold:
            inliers.append(point)

    return np.array(inliers)

def get_random_sample(points_list):
    """Get a list of three random points from a given list"""
    sample_indices = random.sample(range(len(points_list)), 3)
    return np.array([points_list[i] for i in sample_indices])

def calculate_distance_from_center(x_values, y_values, center_x, center_y):
    """Calculate the distance of each 2D point from the circle's center"""
    return np.sqrt((x_values - center_x)**2 + (y_values - center_y)**2)

def algebraic_distance(center, x_values, y_values):
    """Calculate the algebraic distance between points and the mean circle centered at the given center"""
    distances = calculate_distance_from_center(x_values, y_values, *center)
    return distances - distances.mean()

def fit_circle_via_least_squares(x_mean, y_mean, points):
    """Fit a circle using least squares optimization"""
    x_coordinates = points[:, 0]
    y_coordinates = points[:, 1]
    initial_center = x_mean, y_mean
    estimated_center, _ = optimize.leastsq(algebraic_distance, initial_center, (x_coordinates, y_coordinates))

    center_x, center_y = estimated_center
```

Figure 6

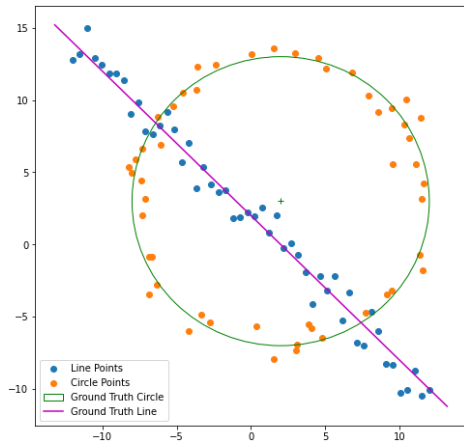


Figure 7 - Original image

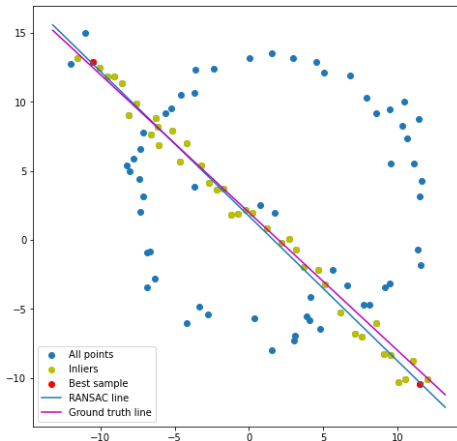


Figure 8- RANSAC Line Fitting

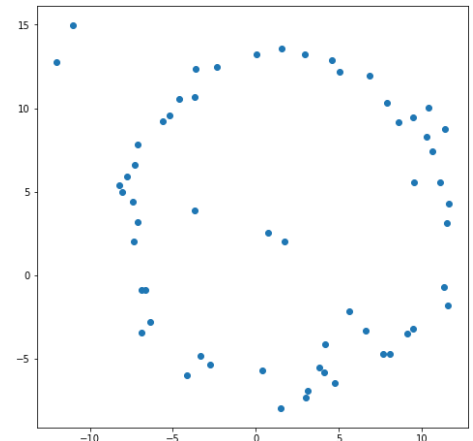


Figure 9 - Detected Outliers and Fitted Circle Visualization

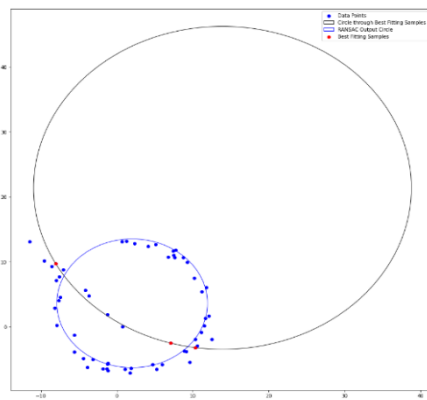


Figure 10- RANSAC Circle Fitting Function

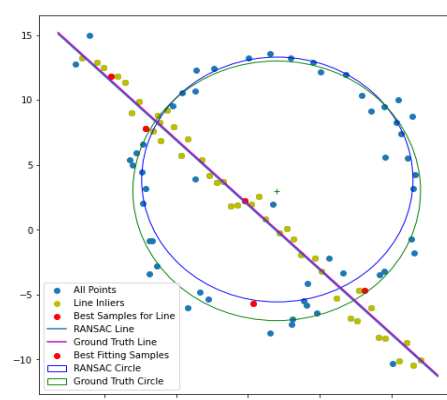


Figure 11 - Revised Plotting Code

- Fitting a circle first on predominantly linear data can lead to poor results because the circle will try to capture all points, even those that follow a line, resulting in a large radius and a poorly positioned center. Many line points might be misclassified as outliers, lowering the inlier ratio and weakening RANSAC performance. Additionally, since a circle requires more parameters than a line, fitting it first may lead to overfitting, reducing RANSAC's effectiveness.

### 3.0. Question 03

This task overlays a flag onto an architectural image using homography by first loading both images and selecting four points on the surface for placement. These points are used to compute a homography matrix that aligns the flag's corners with them. The flag is then warped to match the architectural perspective,

and the images are blended using an `alpha` parameter for transparency, resulting in a seamless integration of the flag into the scene.

```
# Function to compute the homography and warp the flag image onto the architectural image
def warp_flag_onto_image(architecture_img, flag_img, architecture_points, flag_points):
    # Compute the homography matrix
    homography_matrix, _ = cv2.findHomography(flag_points, architecture_points)

    # Warp the flag image to match the architecture image
    warped_flag = cv2.warpPerspective(flag_img, homography_matrix, (architecture_img.shape[1], architecture_img.shape[0]))

    return warped_flag

# Function to blend the warped flag and architectural image
def blend_images(architecture_img, warped_flag, alpha=0.7):
    # Blend the two images
    blended_image = cv2.addWeighted(architecture_img, 1, warped_flag, alpha, 0)
    return blended_image
```





Figure 11



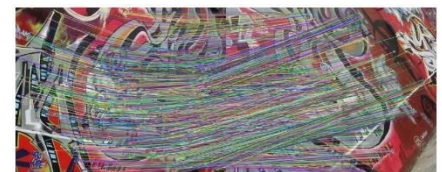
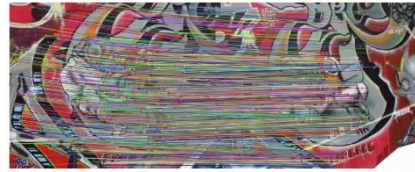
Figure 12



Figure 13

## 4.0. Question – 04

The implementation stitches `img1.ppm` onto `img5.ppm` by first detecting and matching keypoints using the SIFT (Scale-Invariant Feature Transform) algorithm, applying Lowe's ratio test to filter good matches. Next, it employs a custom RANSAC (Random Sample Consensus) method to compute the best homography matrix based on the matched points, ensuring robust estimation. Finally, the homography is used to warp `img1.ppm` and seamlessly stitch it onto `img5.ppm`. This workflow effectively combines feature matching, homography computation, and image warping to create a cohesive stitched image.



```
[[ 4.76241806e-03  2.64303109e-02  2.54729367e-02]
 [-1.46911664e-03  6.40402890e-02  7.62394910e-02]
 [-1.19908222e-03 -7.70266362e-04  1.00000000e+00]]
```

Figure 14- Homography Matrix from RANSAC

```
[[ 6.70443673e+00 -7.22470754e+00 -1.39447473e+00]
 [ 5.29689825e+00 -5.17070305e+00 -2.37703088e+02]
 [ 1.02250666e-02 -1.32862858e-02  1.00000000e+00]]
```

Figure 15 Homography Matrix from RANSAC:

```
def ransac(matched_points):
    maxinliers = 0
    best_H = None
    for i in range(10):
        random_points = random_sample(matched_points)
        homography = calculateHomography(random_points)
        num_inliers = 0
        for i in range(len(matched_points)):
            d = loss(matched_points[i], homography)
            if d < 3:
                num_inliers += 1
        if num_inliers > maxinliers:
            maxinliers = num_inliers
            best_H = homography
    return best_H
```

```
def calculateHomography(correspondences):
    temp_list = []
    for points in correspondences:
        p1 = np.matrix([points.item(0), points.item(1), 1]) * (x1,y1)
        p2 = np.matrix([points.item(2), points.item(3), 1]) * (x2,y2)
        a2 = [0, 0, 0, p2.item(0), p2.item(2) * p1.item(1), p2.item(2) * p1.item(2),
              p2.item(1) * p1.item(0), p2.item(1) * p1.item(1), p2.item(1) * p1.item(2)]
        a1 = [-p2.item(2) * p1.item(0), p2.item(2) * p1.item(1), p2.item(2) * p1.item(2), 0, 0, 0,
              p2.item(0) * p1.item(0), p2.item(0) * p1.item(1), p2.item(0) * p1.item(2)]
        temp_list.append(a1)
        temp_list.append(a2)
    assemble_matrix = np.matrix(temp_list)
    #svd composition
    u, s, v = np.linalg.svd(assemble_matrix)
    #reshape the min singular value into a 3 by 3 matrix
    h = np.reshape(v[8], (3, 3))
    #normalize
    h = (1/h.item(0)) * h
    return h
```

Figure 16- Output