### Image Processing Homework 6

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#### 1

Implement a segmentation algorithm using either K-means clustering or a Gaussian Mixture Model (choose one, and do not use functions from external libraries).

Test the algorithm on images contaminated with noise (e.g., 0.1% salt-and-pepper noise):

- (1) Test binary segmentation and compare the results of your implemented algorithm with those obtained using a thresholding method (such as Otsu's method or based on maximum entropy).
- (2) Test multi-class segmentation (please decide the number of segmentation label classes yourself, with at least three classes).
- (3)Discuss why the segmentation results are inaccurate for noisy images and what methods might yield better segmentation results (note: you are not required to implement these methods, just discuss them).

#### Solution:

For *GMMEM*, the code in gmmem.py depends largely on the slides despite some modifications, and thus shown in the Appendix. The code for *K-means* is in kmeansimg.py is shown below.

```
from PIL import Image
1
     import numpy as np
2
3
     import matplotlib.pyplot as plt
     def kmeans(img, k, max_iter=100, tol=1e-5):
4
5
        Use k-means algorithm to segment the image.
6
7
        # Initialize the centroids by selecting k random pixels from the image
8
        np.random.seed(7) # for reproducibility
9
        centroids = img.ravel()[np.random.choice(img.size, k, replace=False)]
10
11
        height, width = img.shape
12
        labels = np.zeros((height, width), dtype=np.int_)
13
14
        m = 0 # Number of iterations
15
        for _ in range(max_iter):
16
            # Update the labels
17
```

```
distances = (img[..., None] - centroids)**2 # broadcasting to subtract centroids from all
18
                pixels
            labels = np.argmin(distances, axis=2) # find the index of the closest centroid
19
20
            # Update the centroids
21
22
            centroids_new = np.array(
                [img[labels == i].mean() if np.any(labels == i) else 0 for i in range(k)]) # avoid empty
23
                    clusters
24
            # Check the convergence
25
            if np.sqrt(np.sum((centroids_new - centroids)**2)) < tol:</pre>
26
                print('Kmeans Converged! Number of iteration: ', m)
27
28
                break
29
            else:
                centroids = centroids_new
30
                m += 1
31
32
        new_img = np.zeros((height, width), dtype=np.uint8)
33
        for i in range(k):
34
            new_img[labels == i] = centroids[i]
35
36
37
        return labels, centroids, new_img
```

For K-means, we first initialize the centroids by randomly selecting k pixels from the image. Then we update the labels by assigning each pixel to the closest centroid. We then update the centroids by calculating the mean of the pixels in each cluster. The procedure is repeated until the centroids converge or the maximum number of iterations is reached.

For GMMEM, we first initialize the parameters  $\pi_k$  with equal weights,  $\mu_k$  as the centroids obtained from K-means, and  $\sigma_k^2$  as the global variance of the image. Then we perform E-step and M-step alternatively until the parameters converge or the maximum number of iterations is reached.

#### Result:

We test the K-means and GMMEM segmentation on an image polluted with salt-and-pepper noise, and the result is shown in Figure 1.

#### Analysis:

- (1) For binary segmentation, I find that the segmentation result of *K-means* is similar to that of Otsu's method, whereas the performance of *GMMEM* is not as optimal as the other two, because it tends to segment the main part of the heart as a single class, which deviates from our desired outcome. Also, running *GMMEM* is rather slow. I relaxed the conditions for covergence to speed up the process, and the algorithm converged after around 25 iterations.
- (2) In multi-class segmentation, the performance of GMMEM surpasses that of K-means. The processed image is much closer to the original, but there is still room for improvement due to the presence of

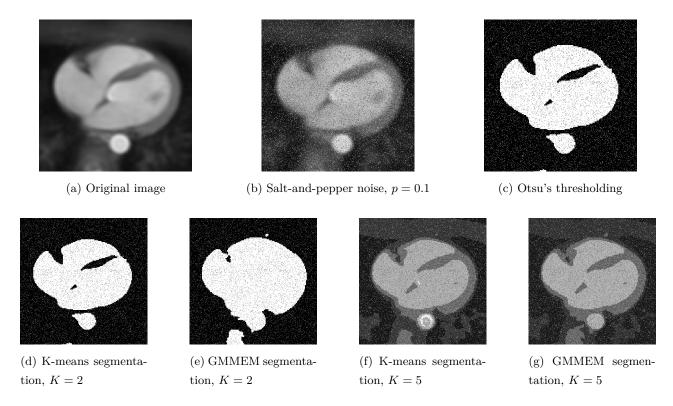


Figure 1: Heart image segmentation

unremoved noise.

(3) The inaccuracy in segmentation results for noisy images stems from our algorithm's inability to effectively recognize and handle noise. I used salt-and-pepper noise, which is sharp and sudden disturbances in the image. This type of noise can significantly alter the pixel intensity distribution, making traditional segmentation algorithms less effective, as they often rely on these distributions.

Before segmentation, using smoothing methods that are less sensitive to outliers such as the median filter can help in reducing the effect of salt-and-pepper noise. Adaptive thresholding may also be applied to improve the segmentation results, as it can automatically adjust the threshold over different parts of the image. Additionally, pre-processing techniques like morphological operations (erosion and dilation) can help in reducing the noise.

#### $\mathbf{2}$

Use morphological operations learned in the course to implement hole filling and removal of isolated points in binary images.

You should code the morphological operations yourself and not use library functions.

#### Solution:

The morphological methods are implemented in morph.py, and is shown below.

```
2
     import numpy as np
3
     import matplotlib.pyplot as plt
4
     def dilation(image, kernel):
5
        k_h, k_w = kernel.shape
6
7
        i_h, i_w = image.shape
8
        # calculate padding size
9
        pad_height = k_h // 2
10
        pad_width = k_w // 2
11
12
        # padding image
13
        padded_image = np.pad(image, ((pad_height, pad_height), (pad_width, pad_width)), mode='constant')
14
        dilated_image = np.zeros_like(image)
15
        for i in range(i_h):
16
            for j in range(i_w):
17
18
               # the neighborhood of the current pixel
               neighborhood = padded_image[i:i+k_h, j:j+k_w]
19
               # apply dilation operation
20
               dilated_image[i, j] = np.max(neighborhood * kernel)
21
22
23
        return dilated_image
24
25
     def erosion(image, kernel):
26
        k_h, k_w = kernel.shape
        i_h, i_w = image.shape
27
28
        # calculate padding size
29
30
        pad_height = k_h // 2
        pad_width = k_w // 2
31
32
        # padding image
33
34
        padded_image = np.pad(image, ((pad_height, pad_height), (pad_width, pad_width)), mode='constant')
        eroded_image = np.zeros_like(image)
35
        for i in range(i_h):
36
            for j in range(i_w):
37
38
               # the neighborhood of the current pixel
39
               neighborhood = padded_image[i:i+k_h, j:j+k_w]
               # apply erosion operation
40
               eroded_image[i, j] = np.min(neighborhood[kernel == 1])
41
42
43
        return eroded_image
44
45
     def opening(image, kernel1, kernel2):
46
        return dilation(erosion(image, kernel1), kernel2)
47
     def closing(image, kernel1, kernel2):
48
        return erosion(dilation(image, kernel1), kernel2)
49
```

For dilation, we construct a kernel with odd size and apply it to every pixel in the image. The kernel is a matrix with 0 and 1, and the size of the kernel is the size of the neighborhood. To avoid the boundary problem, we pad the image with 0 ahead of time. The kernel is applied to the neighborhood of the current pixel, and the intensity is set to white if there is at least one white pixel in the neighborhood overlapping with the "1"s in the kernel. This is equivalent to obtain the maximum intensity where the kernel overlaps with the neighborhood in binary images.

For erosion, the procedure is similar, with the only difference that the intensity is set to white only when all the pixels in the neighborhood overlapping with "1"s in the kernel are white. This is equivalent to obtain the minimum intensity as described above.

Opening is the erosion followed by dilation, and closing is the dilation followed by erosion.

#### Result:

We test the morphological operations on the given binary image and the result is shown in Figure 2.

## ZMIC@FDU ZMIC@FDU

(a) Original image

(b) Image with noise

# ZMIC@FDU ZMIC@FDU

(c) Opening with kernel  $5 \times 5$  and  $9 \times 9$ 

(d) Closing with kernel  $5 \times 5$  and  $9 \times 9$ 

Figure 2: Morphological operations

#### Analysis:

The morphological operations are effective in removing noise and filling holes in the image. After some trials, I found that dilation operation could remove the noise points and lines, while erosion operation could fill the holes in the image. This is aligned with theoretical analysis since our image is black in the foreground and white in the background. In addition, the kernel sizes for both operations have to be larger than  $5 \times 5$  to achieve desired results.

Then we can combine the two operations to obtain better results. The second line above shows the result of opening and closing. I find that both operations with kernel  $5 \times 5$  and thereafter  $9 \times 9$  can reach a desired outcome. We can notice that the left result (opening) proves better, as there are still some gaps in the letter "D" in the right result. The reason is probably that the noises are far from the letters, but some holes are closer to the background. If we first remove the noises by dilation, some letter boundary may be removed as well, and it is difficult to recover by erosion, as shown in the right result. Also due to padding, there is a black frame around the image. However, first filling the holes by erosion will not much affect the letters, and then dilation to remove the noises will give a nice result.

#### A Code for GMMEM

```
from PIL import Image
2
     import numpy as np
3
     import matplotlib.pyplot as plt
     from kmeansimg import kmeans
4
5
6
7
     class Gmmem():
        def __init__(self, img, k, paras):
8
9
            self.img = img # Image to be segmented
            self.num = k # Number of classes
10
            self.shape = img.shape # Shape of the image
11
12
            # Parameters for the GMM, including the weights(\pi_k), means(\mu) and variances(\sigma^2)
13
            self.paras = paras
14
        def phi(self, mean, var, x):
15
            # Normal probability density function
16
17
            return np.exp(-(x - mean) ** 2 / (2 * var)) / np.sqrt(2 * np.pi * var)
18
19
        def get_prob(self, x):
            # Calculate the posterior probability for each class and return the class with the highest
20
                probability
            prob = np.array([self.phi(self.paras[1][i], self.paras[2][i], x) for i in range(self.num)])
21
            posterior = (prob * self.paras[0]) / \
22
                (np.sum(prob * self.paras[0]) + 1e-8) # avoid zero division
23
24
            return int(np.argmax(posterior))
25
        def inference(self):
26
            # Perform inference on the image, and return the mask with the same shape as the image
27
28
            mask = np.array([self.get_prob(i) for i in self.img.reshape(-1)]
                          ).reshape(self.shape[0], self.shape[1])
29
            return mask
30
31
32
        def segmentation(self, max_iter, tol):
            # Perform segmentation on the image, alternatively perform E-step and M-step
33
            m = 0 # Number of iterations
34
            while True:
35
               P = self.E_step() # E-step: Calculate posterior probabilities
36
               # M-step: Update model parameters
37
               new_paras = self.M_update_paras(P)
38
               if (abs(new_paras - self.paras) < tol).all():</pre>
39
                   print('gmm Converged! Number of iteration: ', m)
40
                   break
41
               elif m > max_iter:
42
43
                   break
               else:
44
```

```
45
                   self.paras = new_paras
46
                   m += 1
            mask = self.inference()
47
48
            return mask
49
50
        def P_cal(self, x):
            # Calculate Q/posterior for every pixel
51
            prob = np.array([self.phi(self.paras[1][i], self.paras[2][i], x)
52
                          for i in range(self.num)])
53
            posterior = (prob * self.paras[0]) / \
54
                (np.sum(prob * self.paras[0]) + 1e-8) # avoid zero division
55
            return posterior
56
57
        def E_step(self):
58
            # Calculate the posterior probabilities for each pixel in the image
59
            P = np.stack([self.P_cal(i)
60
61
                      for i in self.img.reshape(-1)], 0).T # (k, h*w)
            return P
62
63
        def M_update_paras(self, P):
64
65
            # Update the model parameters based on the posterior probabilities
            new_paras = np.zeros(self.paras.shape)
66
            new_paras[1] = np.array(
67
68
                [np.sum(P[i] * self.img.reshape(-1)) / np.sum(P[i]) for i in range(self.num)])
69
            new_paras[2] = np.array([np.sum(P[i] * (self.img.reshape(-1) - new_paras[1][i]) ** 2) / np.
                sum(P[i])
                                 for i in range(self.num)])
70
            new_paras[0] = np.sum(P, -1) / np.sum(P)
71
72
            new_paras[2][new_paras[2] < 1] = np.random.randint(1, 10, 1) # Avoid zero variance
            return new_paras
73
74
        def create_newimg(self, max_iter=50, tol=1):
75
76
            # Create a new image based on the labels
            labels = self.segmentation(max_iter, tol)
77
            new_img = np.zeros(self.shape, dtype=np.uint8)
78
            for i in range(self.num):
79
80
               new_img[labels == i] = self.paras[1][i]
81
            return new_img
82
     def initialize_paras(img, k):
83
84
        paras = np.zeros((3, k))
        paras[0] = np.ones(k) / k # equal weights
85
        labels, centroids, _ = kmeans(img, k)
86
        paras[1] = centroids
87
88
        paras[2] = np.array([np.var(img[labels == i]) for i in range(k)])
89
        return paras
```