Machine Learning Homework 6 Kernel K-means and Spectral Clustering

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- o a. code with detailed explanations (40%)
 - Part1 (15%)
 - ** whole project set gammac = 1/(255*255), gammas = 1/(100*100)
 - (a). Kernel K-means

```
50  def KernelFunction(pixel, coord):
51     s = np.exp(-gamma_s * cdist(coord, coord, 'sqeuclidean'))
52     c = np.exp(-gamma_c * cdist(pixel, pixel, 'sqeuclidean'))
53
54     return s * c
```

Compute kernel function using RBF kernel on both spatial information and color information and multiply them as complex kernel.

Do the computation of formula in slide p.22

$$\begin{aligned} \left\| \phi(x_j) - \mu_k^{\phi} \right\| &= \left\| \phi(x_j) - \frac{1}{|C_k|} \sum_{n=1}^N \alpha_{kn} \phi(x_n) \right\| \\ &= \mathbf{k}(x_j, x_j) - \frac{2}{|C_k|} \sum_n \alpha_{kn} \mathbf{k}(x_j, x_n) + \frac{1}{|C_k|^2} \sum_p \sum_q \alpha_{kp} \alpha_{kq} \mathbf{k}(x_p, x_q) \end{aligned}$$

```
def KernelKMeans(imgname, savepath, pixel, coord):
   init_method = ['random', 'modK']
   for method in init_method:
         gif = []
          FOLDER = f'{K_cluster}_cluster_{method}'
         save_method_path = f'{savepath}/{FOLDER}'
if not os.path.exists(save_method_path):
              os.makedirs(save_method_path)
         cluster = initial(pixel, method)
kernel = KernelFunction(pixel, coord)
         iteration = 0
         error = -100000
         prev_error = -100000
while(iteration < EPOCH):</pre>
            iteration += 1
              print("iteration = {}".format(iteration))
              prev_cluster = cluster
               img = Visualization(imgname, save_method_path, iteration, cluster, method)
              gif.append(img)
               cluster = clustering(pixel, kernel, cluster)
              error = ComputeError(cluster, prev_cluster)
               print("error = {}".format(error))
               if error == prev_error:
              prev_error = error
```

Summarize:

step1: Get parameters using sys.argv

step2: Calculate kernel using spatial information and color information

step3: Use different initialized method to cluster each pixel

step4: Calculate distance between pixel and center

step5: Update cluster

(b). Spectral Clustering

```
def KernelFunction(pixel, coord):
    # compute spatial information RBF kernel

s = np.exp(-gamma_s * cdist(coord, coord, 'sqeuclidean'))

c = np.exp(-gamma_c * cdist(pixel, pixel, 'sqeuclidean'))

return s * c
```

Compute kernel regarding spatial information and color information

```
# update clustering results using new centriods
def clustering(E, centroids):

cluster = np.zeros(scale * scale, dtype = int)
for i in range(scale * scale):
    distance = np.zeros(K_cluster, dtype = np.float32)
    for j in range(K_cluster):
        distance[j] = np.sum(np.absolute(E[i] - centroids[j]))
    cluster[i] = np.argmin(distance)
    return cluster
```

Do clustering of each pixel

```
# update centroids using

def UpdateCentroids(E, centroids, cluster):
    centroids = np.zeros(centroids.shape, dtype=np.float64)

tmp = np.zeros(K_cluster, dtype=np.int32)

for i in range(scale * scale):
    centroids[cluster[i]] += E[i]

tmp[cluster[i]] += 1

for i in range(K_cluster):
    if tmp[i] == 0:
    tmp[i] == 0:
    tmp[i] = 1

centroids[i] /= tmp[i]

return centroids
```

Update the new centroids according to the clustering results

```
def KMeans(imgname, E, cut, init_method):
    centroids, cluster = initial(E, init_method)
    Visualization(imgname, cluster, iteration, cut)
    prev_error = -100000
    error = -100000
    error = -100000
    if = []
    iteration = 0
    while iteration < EPOCH:
        iteration += 1
        prev_cluster = cluster
        cluster = clustering(E, centroids)
        centroids = UpdateCentroids(E, centroids, cluster)
        img = Visualization(imgname, cluster, iteration, cut)
        gif.append(img)
        error = ComputeError(cluster, prev_cluster)
    print(f'Iter: {iteration}: {error}')
    if error == prev_error:
        break
    prev_error = error
    if K_cluster == 2:
        EigenSpace(E, cluster, cut)</pre>
```

I run 15 epoch to do the iteration, if error == prev error than break the loop.

Summarize:

step1: Get parameters using sys.argv

step2: Calculate kernel using spatial information and color information

step3: Calculate Laplacian L

step4: Calculate the first k eigenvectors of L

step5: Use eigenvectors to do K-means

- Part2 (5%)
- (a) Kernel K-means

Running command: python file.py imgname k_cluster, if want to have more cluster, then change the 'k_cluster' parameter.

EX: python kernelkmeans.py image1.png 3, python spectralclustering.py image1.png 4

(b) Spectural Clustering

Running command: python file.py imgname k_cluster initialization cut, if want to have more cluster, then change the 'k cluster' parameter.

EX: python spectralclustering.py image1.png 3 kmeans++ ratio, python spectralclustering.py image2.png 3 random normalized

- Part3 (10%)
- (a) Kernel K-means

```
def initial(pixel, initial_method):
    if initial_method == 'random':
        classification = np.random.randint(K_cluster, size = pixel.shape[0])
    elif initial_method == 'modK':
        classification = []
    for i in range(data.shape[0]):
        classification.append(i % K_cluster)
    classification = np.asarray(classification)
    return classification
```

I use two initialization method: random and modK to do the cluster initialization.

(b) Spectral Clustering

```
def initial(E, init_method):
    cluster = np.random.randint(0, K_cluster, scale * scale)
# initialization using random
if init_method == 'random':
high = E.max(axis=0)
low = E.min(axis=0)
diff = high - low
centroids = np.random.rand(K_cluster, K_cluster)
for i in range(K_cluster):
    centroids[:, i] *= diff[i]
    centroids[:, i] += low[i]
# initialization using kmeans++
elif init_method == 'kmeans++':
    centroids = [E[np.random.choice(range(scale * scale)), :]]
# find #K_cluster centroids
for i in range(K_cluster - 1):
    dist = cdist(E, centroids, 'euclidean').min(axis=1)
    prob = dist / np.sum(dist)
    centroids = np.array(centroids)

return centroids, np.array(cluster)
```

In spectral clustering, I used two initial method: random and kmeans++ to do initialization.

• Part4 (10%)

```
# use different cut

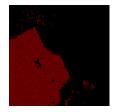
| 144 | def CUT(W, D, cut):
| 145 | if cut == 'normalized':
| 146 | D_Square = np.diag(np.power(D, -0.5))
| 147 | L = np.identity(scale * scale) - D_Square @ W @ D_Square
| 148 | EigenValue, EigenVector = np.linalg.eig(L)
| 149 | idx = np.argsort(EigenValue)[1: K_cluster+1]
| 150 | U = EigenVector[:, idx].real.astype(np.float32)
| 151 | T = np.zeros(U.shape, dtype = np.float64)
| 152 | for i in range(scale * scale):
| 153 | T[i] = U[i] / np.sqrt(np.sum(U[i] ** 2))
| 154 |
| 155 | return T |
| 156 | elif cut == 'ratio':
| 157 | L = D - W |
| 158 | EigenValue, EigenVector = np.linalg.eig(L)
| idx = np.argsort(EigenValue)[1: K_cluster+1]
| 160 | U = EigenVector[:, idx].real.astype(np.float32)
| 161 | return U
```

To do different cut method: normalized cut and ratio cut by calculating eigenvalue and eigenvector of laplacian matrix.

- o b. experiments settings and results (20%) & discussion (30%)
 - Part1 (5%) & (5%)
 - (a) Kernel K-means: cluster number = 2 (initialization: random)
 - 1. Image1







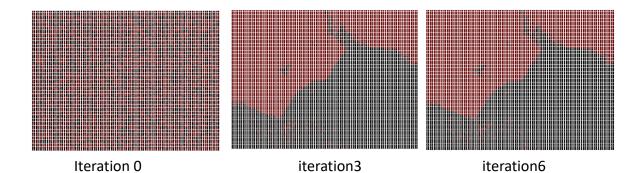
Iteration 0~15

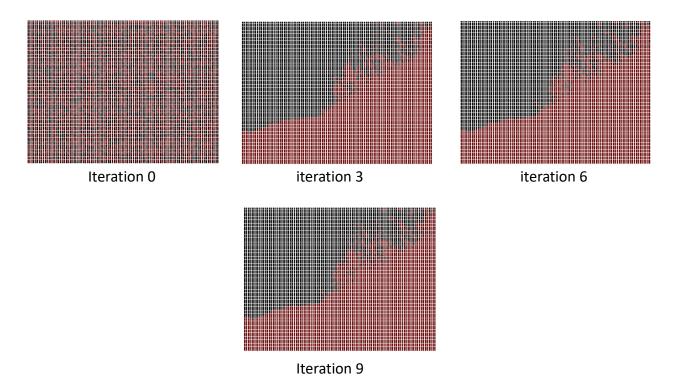


Iteration 0~15

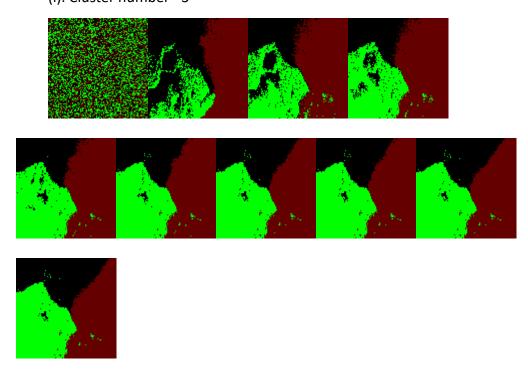
(b) Spectural Clustering : cluster number = 2(initialization: random, cut: normalized)

1. Image1



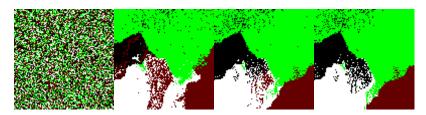


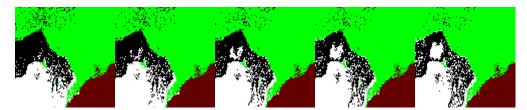
- Part2 (5%) & (5%)
- (a) Kernel K-means (initialization: random)
- 1. Image1
 - (i). Cluster number =3



Iteration (0~15)

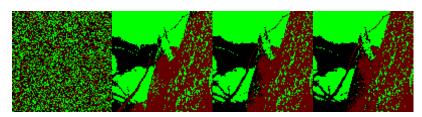
(ii). Cluster number = 4

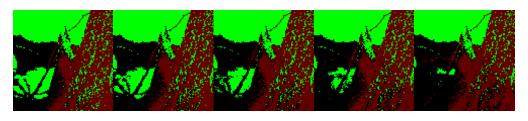


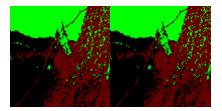




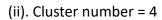
Iteration (0~15)

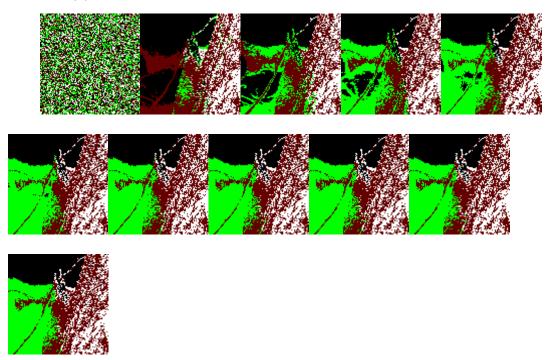






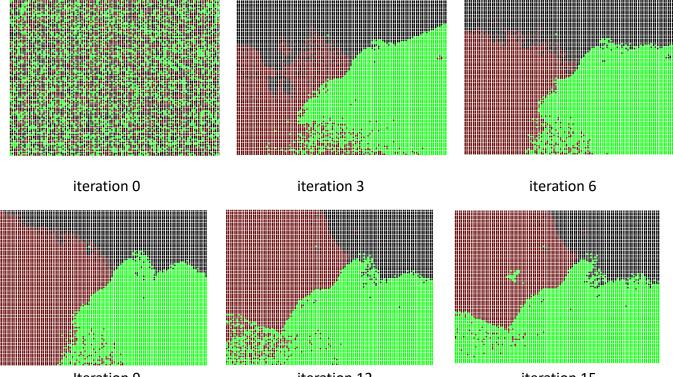
Iteration 0~15



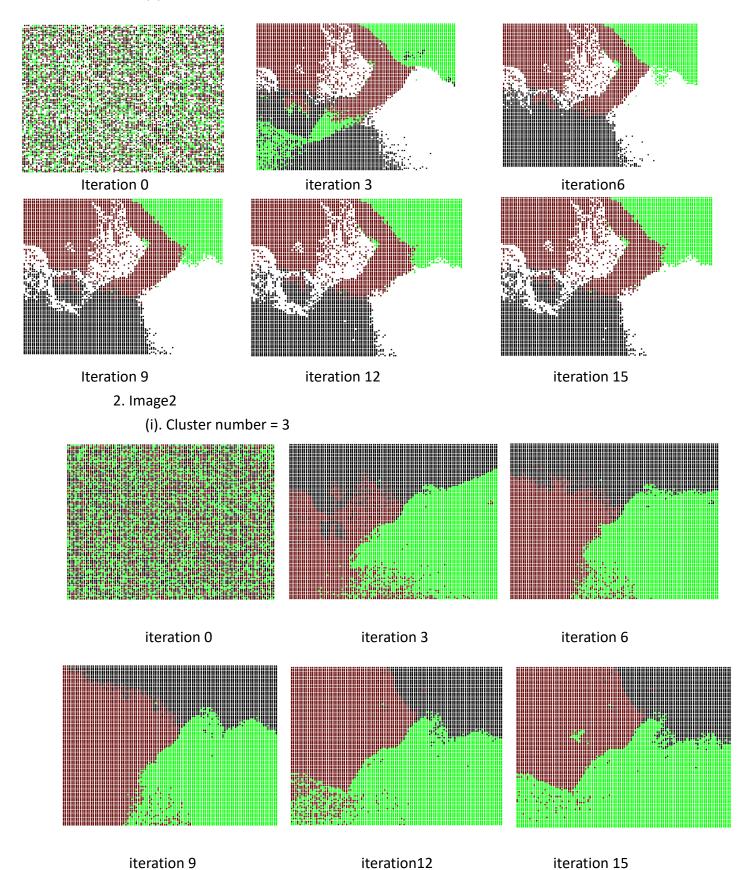


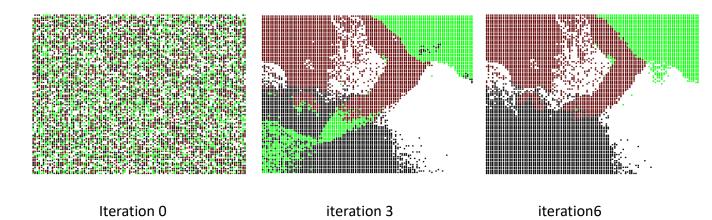
Iteration 0~15

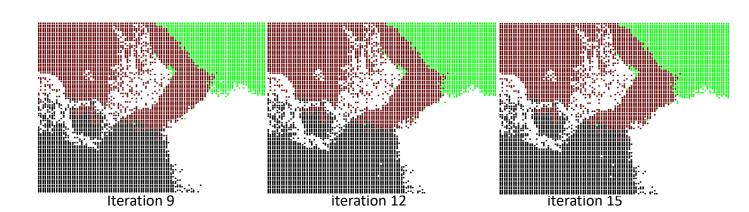
- (b) Spectural Clustering (initialization: random, cut: normalized)
- 1. Image1
 - (i). Cluster number =3



Iteration 9 iteration 12 iteration 15



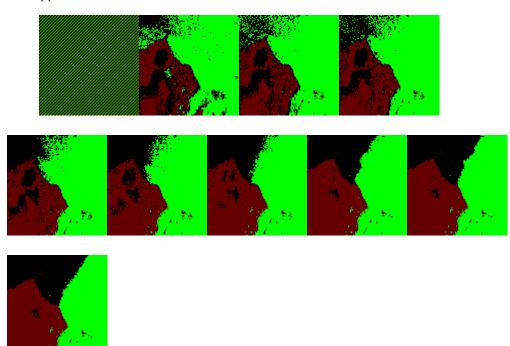




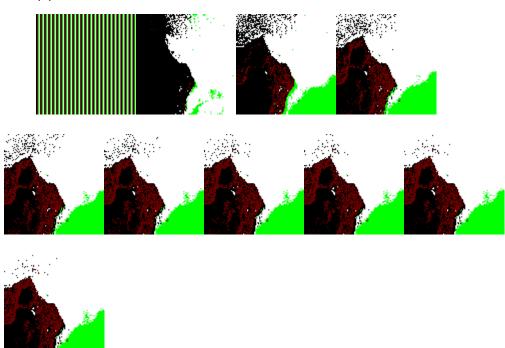
- Part3 (5%) & (10%)
- (a) Kernel K-means

Initialization: mod k (results of random initialization method has shown above.)

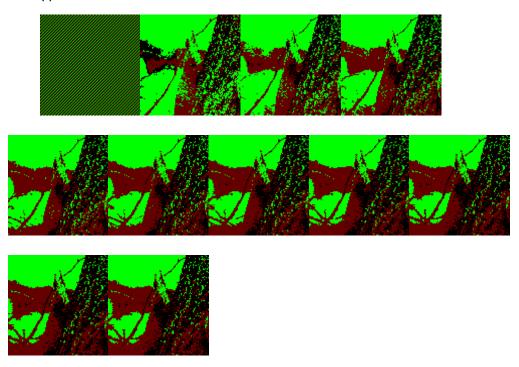
- 1. Image1
 - (i). Cluster number =3



Iteration 0~15

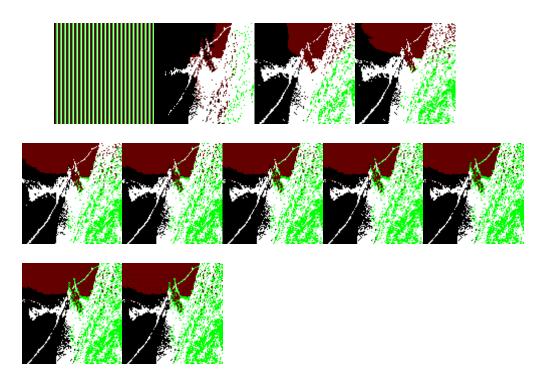


Iteration 0~15



Iteration 0~15

(ii). Cluster number = 4

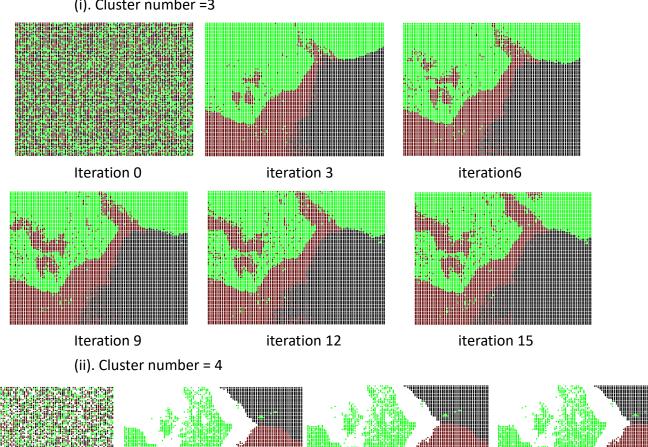


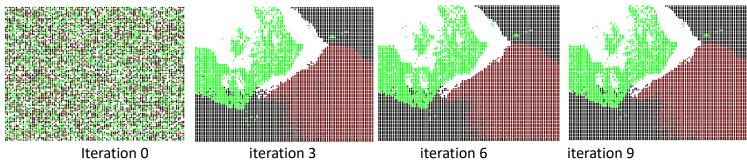
Iteration 0~15

(b) Spectural Clustering Initialization: kmeans++ (results of random initialization method has shown above.)

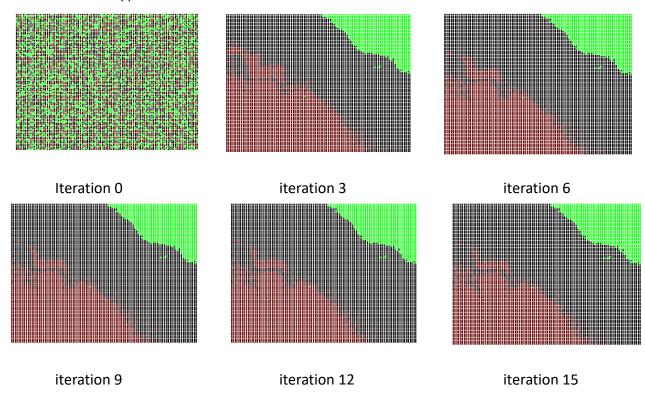
1. Image1

Normalization cut

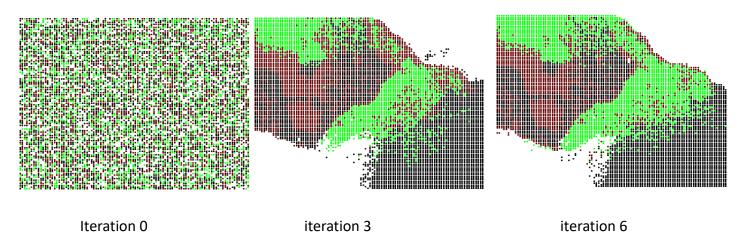


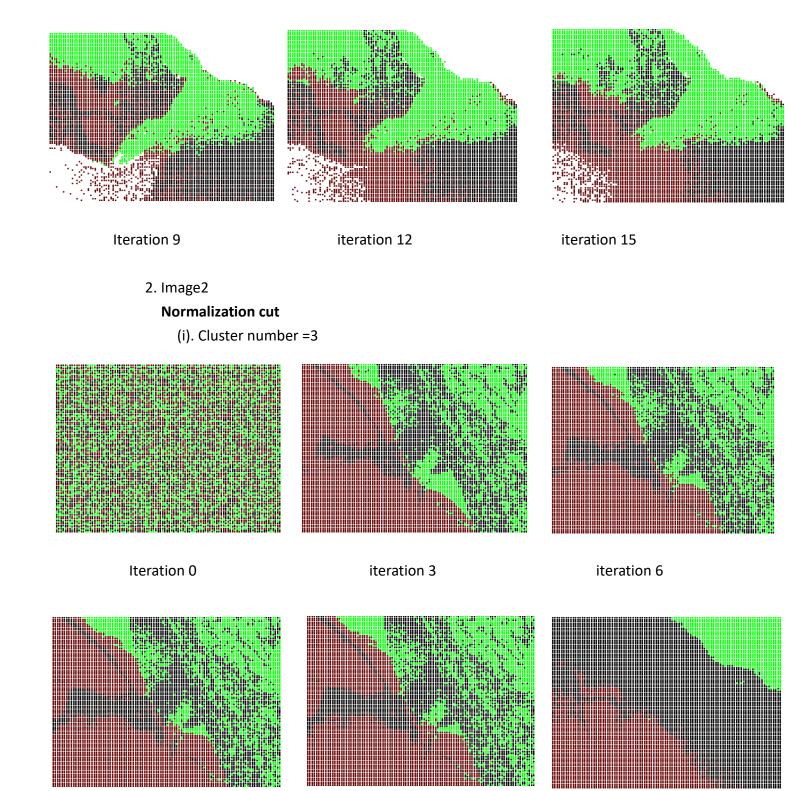


Ratio cut



(ii). Cluster number = 4

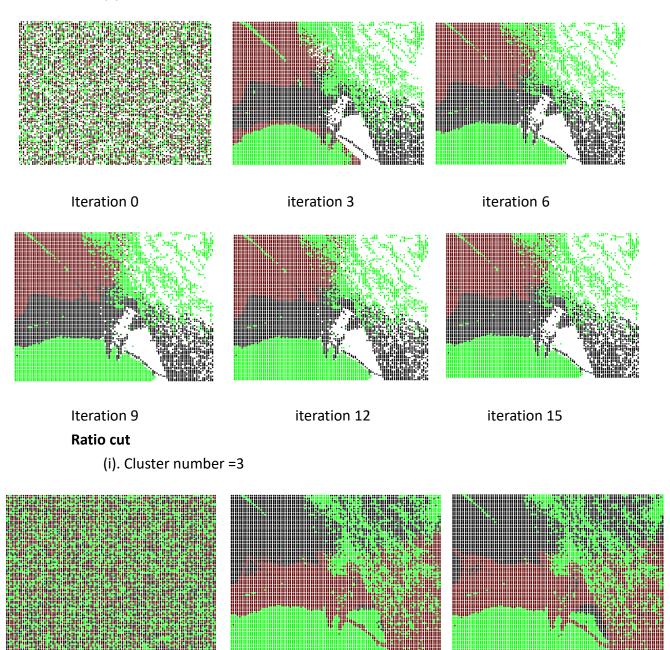




Iteration 9 iteration 12 iteration 15

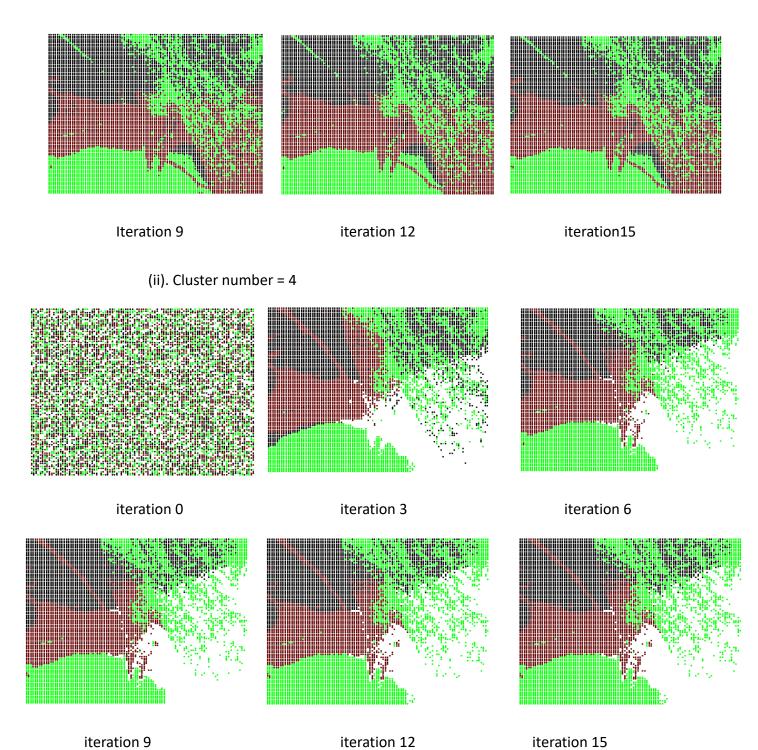
(ii). Cluster number = 4

Iteration 0



iteration 3

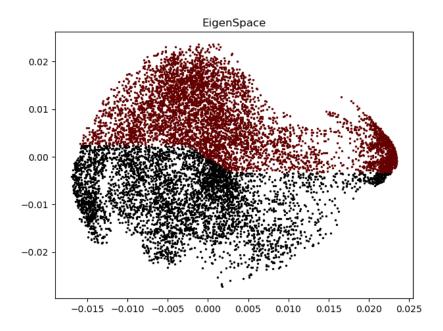
iteration 6



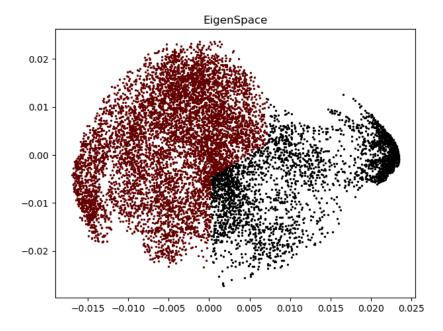
• Part4 (5%) & (10%)

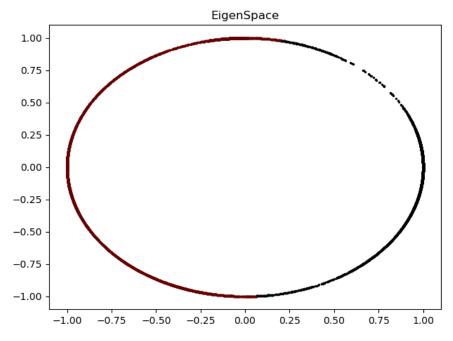
Eigen space of spectral clustering

I. Image1 (k_cluster = 2, Ratio cut, Init: Kmeans++)

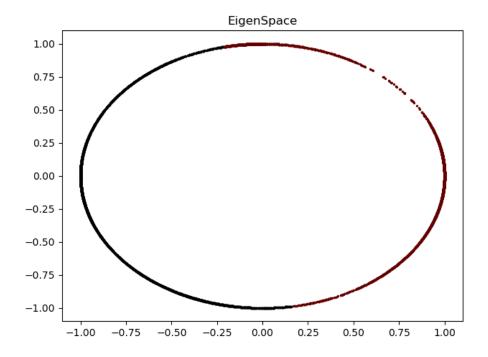


(k_cluster = 2, Ratio cut, Init: random)

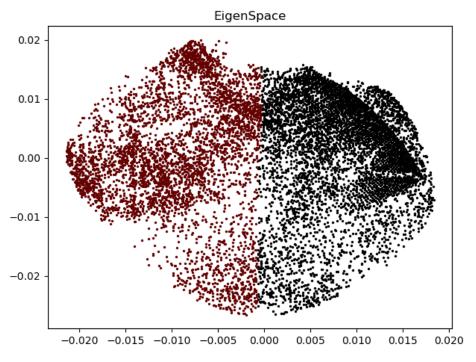




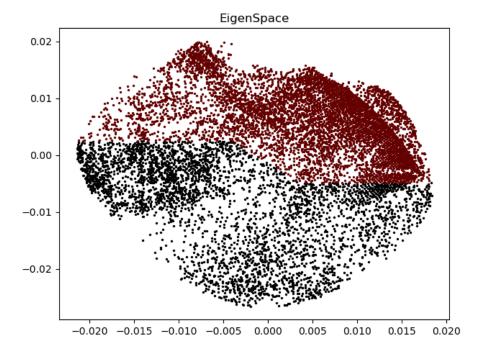
(k_cluster = 2, Normalized cut, Init: random)



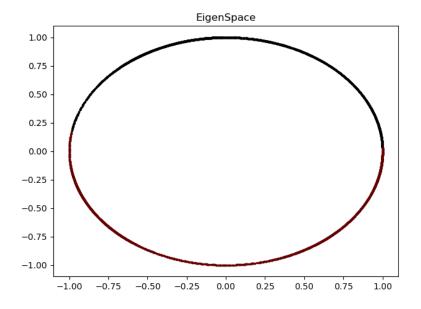
2. Image2 (k_cluster = 2, Ratio cut, Init: Kmeans++)



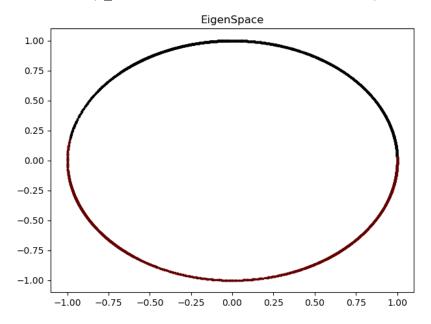
(k_cluster = 2, Ratio cut, Init: random)



(k_cluster = 2, Normalized cut, Init: kmeans++)

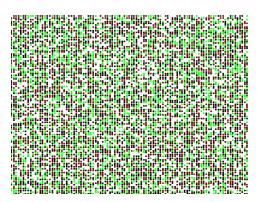


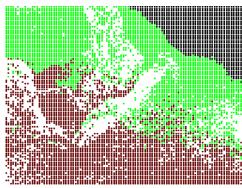
(k_cluster = 2, Normalized cut, Init: random)

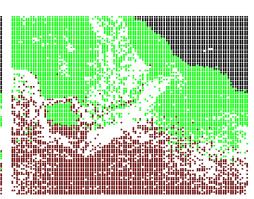


o c. observations and discussion (10%)

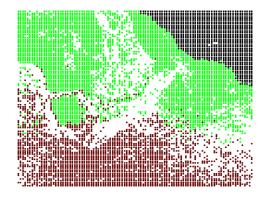
In this lab, I tried to fine tune parameters <code>gamma_c</code> and <code>gamma_s</code>, but it turned out that the results become strange. So I tried to set <code>gamma_c</code> = 1e-5 and <code>gamma_s</code> = 1e-5 to do the experiment on (image1, ratio cut, random initialization), and the results are shown bellow:







Iteration 0 iteration 3 iteration 6



Iteration 9