

# HW6

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## Data(image) Preparation

### Data (image)

```
im = imageio.imread('image1.png').astype('float32')
im2 = imageio.imread('image2.png').astype('float32')
```

In use *imageio* to read image data in both algorithms, which can read image as array (100x100).

### Similarity matrix (W)

```
W1 = rbf_kernel(im,1)
W2 = rbf_kernel(im2,2)
```

We will use similarity matrix in both algorithms for two input images. The matrix is made up by given **RBF kernel**

```
def rbf_kernel(X,image_number):
    sigma_s = -0.02
    sigma_c = -0.02
    W = np.zeros((10000,10000))
    for i in range(10000):
        x1 = i//100
        y1 = i%100
        for j in range(i+1,10000):
            x2 = j//100
            y2 = j%100
            if image_number==1:
                w = np.exp(sigma_s*((x1-x2)**2+(y1-y2)**2))*np.exp(sigma_c*(np.linalg.norm(im[x1,y1]-im[x2,y2])**2))
            else:
                w = np.exp(sigma_s*((x1-x2)**2+(y1-y2)**2))*np.exp(sigma_c*(np.linalg.norm(im2[x1,y1]-im2[x2,y2])**2))
            W[i,j] = W[j,i] = w
    return W
```

which is equivalent to

$$k(x, x') = e^{-\gamma_s \|S(x) - S(x')\|^2} \times e^{-\gamma_c \|C(x) - C(x')\|^2}$$

Where  $S(x)$  is the spatial information and  $C(x)$  is the color information.

# Kernel K-means

## Initial Methods

I tried two different initial methods.

1. **Random assign:** randomly assign data point into class(0~k-1)

```
def init(method,k,image_number):  
    if method == 0:  
        print('Init method: random')  
        return np.random.randint(k,size=10000)
```

2. **K-means++:** from Line 26 – Line 58

In this method, I'll explain in pseudocode followed:

- A. Randomly choose a data point as the first cluster
- B. Calculate distances of each points to the closest clusters
- C. Turn distance into probability and choose next cluster by probability(The farther the higher probability to be choosed)
- D. Repeat B-C till find k cluster.
- E. (from Line 47)Assign each point into their closet cluster.

## Clustering

After initial, I use the kernel data (similarity matrix) and the initial information to cluster data points into k(2~4) clusters.

```
def classify(kernel_data,res_prev,k):  
    res = np.zeros(10000)  
    unique, counts = np.unique(res_prev, return_counts=True)  
    third = third_term(kernel_data,res_prev,k)  
    for j in range(10000):  
        temp = np.zeros(k)  
        for c in range(k):  
            temp[c] += (0-2*second_term(kernel_data,res_prev,j,c)/counts[c]+third[c]/counts[c]**2)  
        res[j] = np.argmin(temp)  
    return res
```

First, calculate the distance from kernel data to the means:

$$\|\phi(x_j) - \mu_k^\phi\| = k(x_j, x_j) - \frac{2}{|C_k|} \sum_n \alpha_{kn} k(x_j, x_n) + \frac{1}{|C_k|^2} \sum_p \sum_q \alpha_{kp} \alpha_{kq} k(x_p, x_q)$$

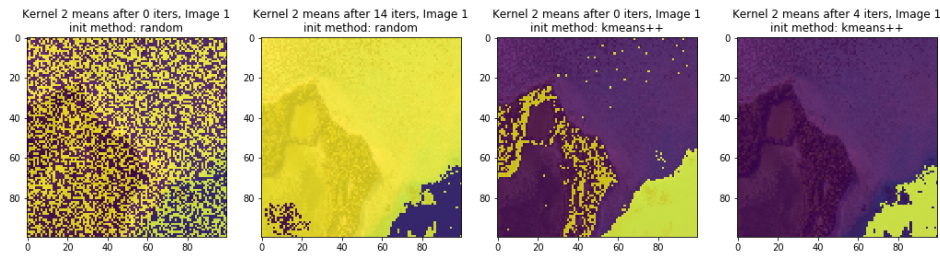
Then cluster each point into closest cluster use *argmin*.

Repeat this step until the difference in clustering result between two iteration converges. (By experiment, I set the stopping threshold  $< 50$ )

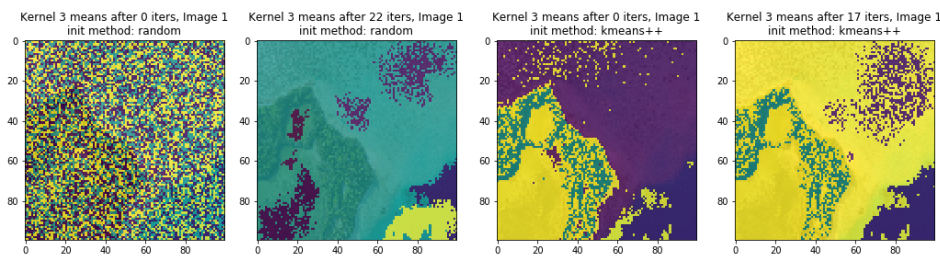
## Results

- Image 1

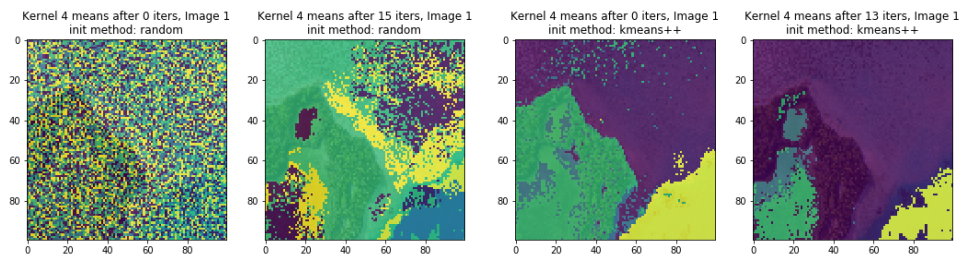
- K=2



- K=3

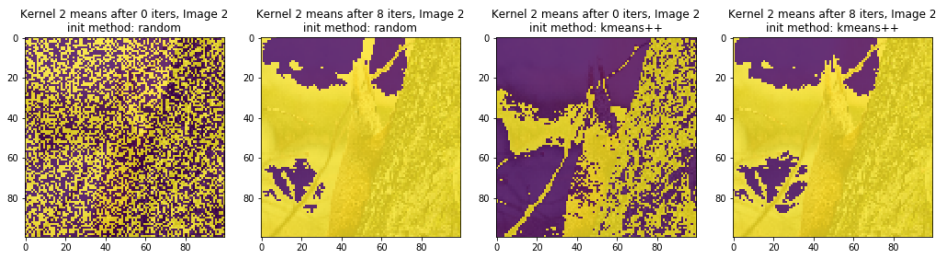


- K=4

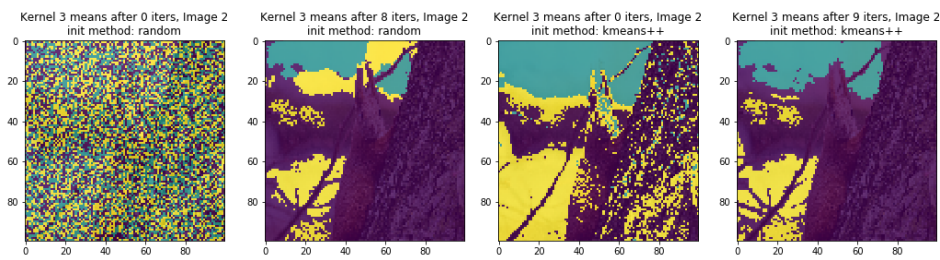


- Image 2

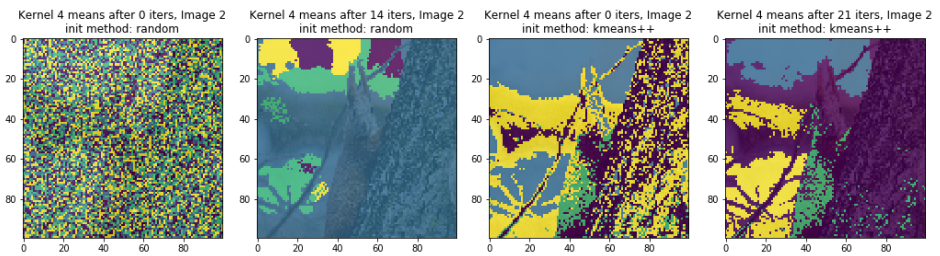
- K=2



- K=3



- K=4



From the result we can see that in both input image, if we use kmeans++ as the initial method, since the starting center is far enough, it needs lesser time to converge and get more precise results.

# Spectral Clustering

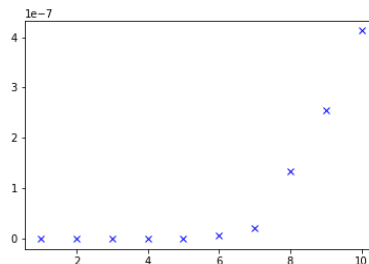
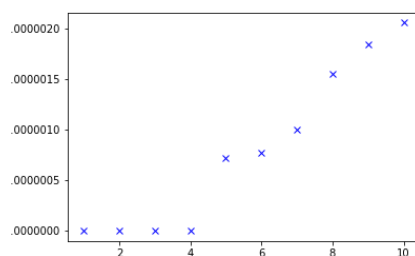
## Ratio-Cut

To approximate the minimum of ratio-cut method, we can use unnormalized Laplacian  $L=D-W$  to approach, where  $D$  is the degree matrix,  $W$  is the similarity matrix.

```
D1 = np.diag(np.sum(W1,axis=1))
D2 = np.diag(np.sum(W2,axis=1))
L1 = D1-W1
L2 = D2-W2
```

Then we find the eigenvalues and eigenvectors of  $L$ , sort eigenvalues to find the top  $k+1$  small eigenvalues and their eigenvectors. Since we use **RBF kernel** here, which cause the fully-connected graph, we start from the 2<sup>nd</sup> smallest eigenvalue and its eigenvector. Let  $U$  be the matrix containing these  $k$  eigenvectors.

```
# Ratio-Cut
eigen_values_1, eigen_vectors_1 = np.linalg.eig(L1)
eigen_values_2, eigen_vectors_2 = np.linalg.eig(L2)
eigen_vectors_1_sort = eigen_vectors_1[:,np.argsort(eigen_values_1)]
eigen_vectors_2_sort = eigen_vectors_2[:,np.argsort(eigen_values_2)]
for k in range(2,5):
    U1 = (eigen_vectors_1_sort[:,1:k+1])
    U2 = (eigen_vectors_2_sort[:,1:k+1])
    k_means(k,U1,1,'ratio cut')
    k_means(k,U2,2,'ratio cut')
```



(Sorted eigen-value of

image1 and image2)

## Normal-Cut

We can use normalized Laplacian  $L_{sym} = D^{-1/2}LD^{-1/2}$  to approximate the minimum of normalized-cut. Then we also start from the 2<sup>nd</sup> smallest eigenvalue and its eigenvector of  $L_{sym}$ , let

$U$  be the matrix of these  $k$  eigenvectors.

```
# Normal-Cut
D1_inv = np.linalg.inv(D1**0.5)
D2_inv = np.linalg.inv(D2**0.5)
L1_sym = np.dot(np.dot(D1_inv,L1),D1_inv)
L2_sym = np.dot(np.dot(D2_inv,L2),D2_inv)
eigen_values_normal_1, eigen_vectors_normal_1 = np.linalg.eig(L1_sym)
eigen_values_normal_2, eigen_vectors_normal_2 = np.linalg.eig(L2_sym)
eigen_vectors_normal_1_sort = eigen_vectors_normal_1[:,np.argsort(eigen_values_normal_1)]
eigen_vectors_normal_2_sort = eigen_vectors_normal_2[:,np.argsort(eigen_values_normal_2)]
for k in range(2,5):
    U1_normal = (eigen_vectors_normal_1_sort[:,1:k+1])
    U2_normal = (eigen_vectors_normal_2_sort[:,1:k+1])
    k_means(k,U1_normal,1,'normal cut')
    k_means(k,U2_normal,2,'normal cut')
```

Then we cluster the vector corresponding to the *rows of  $U$*  into  $k$  clusters with *k-means* algorithm.

## Initial Methods

I tried two different initial methods.

1. **Random assign:** randomly assign  $k$  data points to be the center

```
def init(U,method,k,image_number):
    means = np.zeros((k,k))
    if method == 0:
        print('Init method: random')
        temp = np.random.randint(10000,size=k)
        for i in range(k):
            means[i] = U[temp[i]]
```

2. **K-means++:** from Line 30 – Line 42

In this method, I'll explain in pseudocode followed:

- A. Randomly choose a data point as the first center
- B. Calculate distances of each points to the closest center
- C. Turn distance into probability and choose next center by probability(The farther the higher probability to be choosed)
- D. Repeat B-C till find  $k$  centers.

# K-means

From Line 78 – Line 97, I implement the k-means algorithm:

1. Initial the centers using algorithm introduced aboved.
2. Cluster each point to their closest center (function classify, Line 45 – Line 52)
3. Update center according to the point in the corresponding cluster. (function update, Line 54 – Line 63)
4. Draw the result use matplotlib.

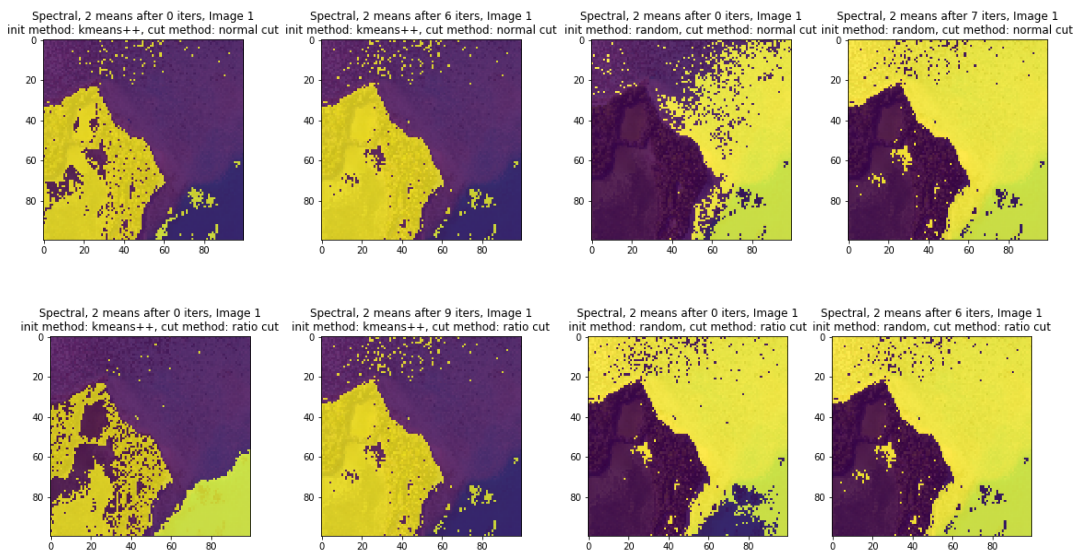
## Results

First line: normal-cut

Second line: ratio-cut

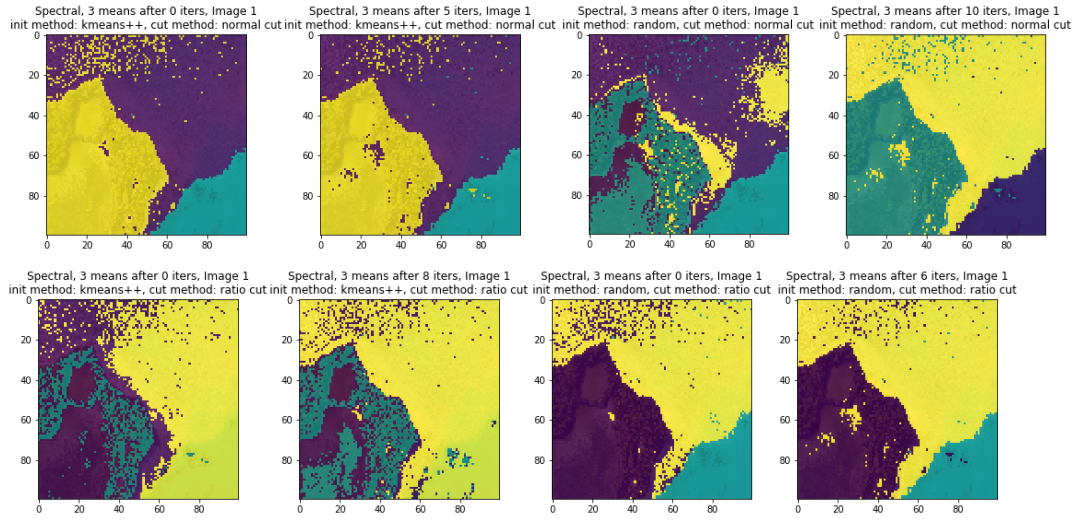
### ● Image 1

#### ■ K=2

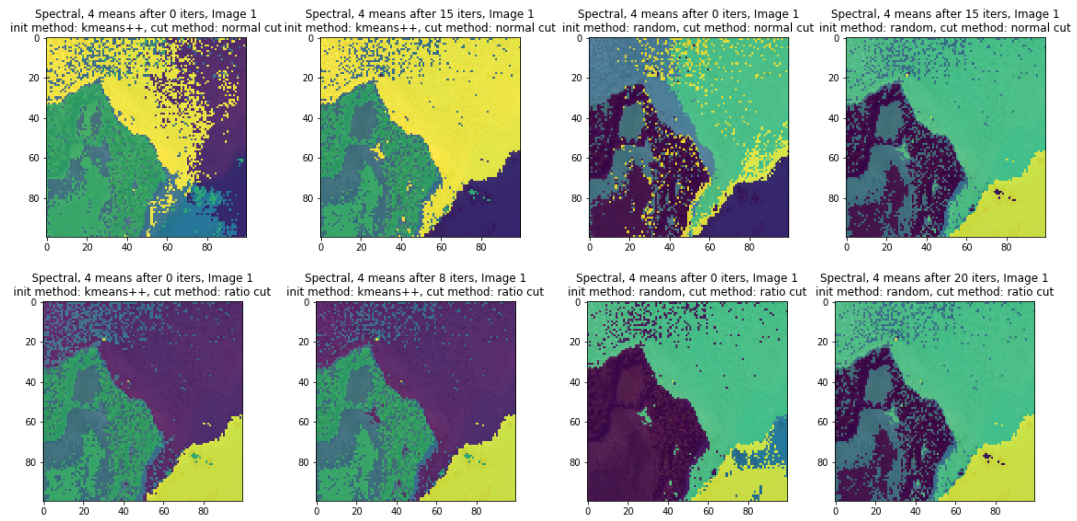




## ■ K=3



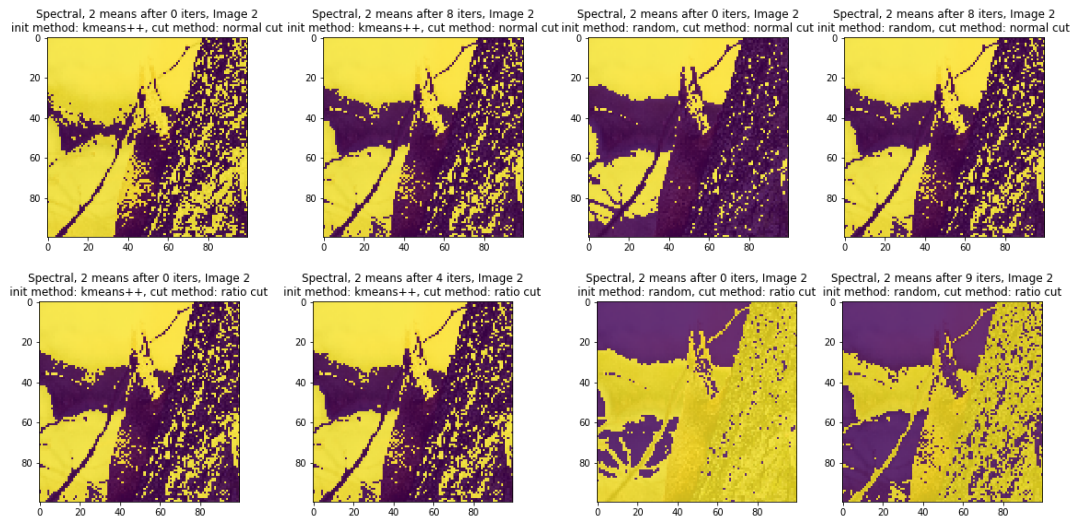
## ■ K=4



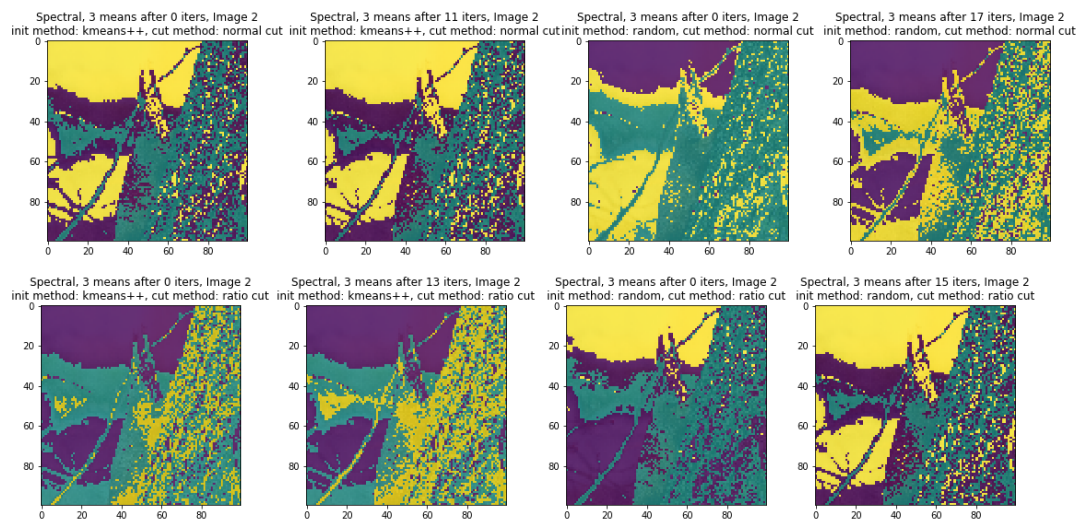


- Image 2

- K=2



- K=3



## ■ K=4

