ML Homework 6 Report

Student Info

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- Code
 - 1. Kernel K-means
 - Common Parts
 - 1. Load Data

```
def readImagesInNpArray():

# Read the image files.
images = [Image.open(filePathOfImage1), Image.open(filePathOfImage2)]

# Convert image to numpy array.
images[0] = np.asarray(images[0])
images[1] = np.asarray(images[1])

return images
```

The objective of the readImagesINpArray function is to load data from the input file, it will be called in the main function.

2. Compute Kernel

```
def computeKernel(image, gammaC):

# Get image shape.

rows, cols, colors = image.shape

# Compute the color distance.

numOfPixels = rows * cols

colorDist = cdist(image.reshape(numOfPixels, colors), image.reshape(numOfPixels, colors), 'sqeuclidean')

# Compute the indices of a grid.

indices = np.indices((rows, cols))

indicesOfRow = indices[0]

indicesOfCol = indices[1]

# Compute the indices vector.

indicesVector = np.hstack((indicesOfRow.reshape(-1, 1), indicesOfCol.reshape(-1, 1)))

# Compute the spatial distance.

spatialDist = cdist(indicesVector, indicesVector, 'sqeuclidean')

# The kernel formula in spec.

return np.multiply(np.exp(-gammaS * spatialDist), np.exp(-gammaC * colorDist))
```

The function is to compute the gram matrix based on the kernel function defined in spec, and the formula is following pic.

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

- Part 1 (2-clusters / randomly init)
 - 1. Init Center

For the init center code of part 1, I just initialize the center randomly according to the number of number of clusters.

2. Init Clusters

```
def initClusters(numOfRows, numOfCols, numOfClusters, kernel, initMode):

# Init centers.

centers = initCenters(numOfRows, numOfCols, numOfClusters, initMode)

# K-means.

numOfPixels = numOfRows * numOfCols

clusters = np.zeros(numOfPixels, dtype=int)

for pixel in range(numOfPixels):

# Compute the distance of every pixel to all centers.

distance = np.zeros(numOfClusters)

for index, center in enumerate(centers):

seqOfCenter = center[0] * numOfRows + center[1]

distance[index] = kernel[pixel, pixel] + kernel[seqOfCenter, seqOfCenter] - 2 * kernel[pixel, seqOfCenter]

# Pick the index of minimum distance as the cluster of the point

clusters[pixel] = np.argmin(distance)

return clusters
```

After get the init center from <u>initcenter</u> function, We will classify all the pixels based on the min distance in feature space (each point to the center).

3. Kernel K-means

```
# Kernel k-means.
currentClusters = clusters.copy()
count = 0
iteration = 100

while True:
    # Compute new clusters.
    numOfPixels = numOfRows * numOfCols
    newClusters = kernelClustering(numOfPixels, numOfClusters, kernel, currentClusters)

# Get the image state.
imageState = getCurrentImageState(numOfRows, numOfCols, newClusters)

imageStates.append(imageState)

if np.linalg.norm((newClusters - currentClusters), ord=2) < 0.001 or count >= iteration:
    break

currentClusters = newClusters.copy()
count += 1
```

After getting the init clusters, we start to perform kernel k-means. In each round, we will perform kernelCluster function to get new clusters and calculate the difference between the currentClusters and newClusters to check if it is already converged.

4. Kernel Clustering

```
# Get number of members in each cluster

# Get number of members in each cluster

numOfMembers = np.array([np.sum(np.where(clusters == c, 1, 0)) for c in range(numOfClusters)])

# Get sum of pairwise kernel distances of each cluster

pairwise = getSumOfPairwiseOistance(numOfPixels, numOfClusters, numOfMembers, kernel, clusters)

newClusters = np.zeros(numOfPixels, dtype=int)

for p in range(numOfPixels):

distance = np.zeros(numOfClusters)

for c in range(numOfClusters):

distance[c] + kernel[p, p] + pairwise[c]

# Get distance from given data point to others in the target cluster

distToOthers = np.sum(kernel[p, :][np.where(clusters == c)])

distance[c] -= 2.8 / numOfMembers[c] * distToOthers

newClusters[p] = np.argmin(distance)
```

We perform the kernelclustering based on the following pic formula. It the end, it will return the new clusters in array form.

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

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- Part 2 (3-clusters / randomly init)
 - 1. Init Center

It is same to the init center code of part 1.

2. Init Clusters

```
def initClusters(numOfRows, numOfCols, numOfClusters, kernel, initMode):

# Init centers.

centers = initCenters(numOfRows, numOfCols, numOfClusters, initMode)

# K-means.

numOfPixels = numOfRows * numOfCols

clusters = np.zeros(numOfPixels, dtype=int)

for pixel in range(numOfPixels):

# Compute the distance of every pixel to all centers.

distance = np.zeros(numOfClusters)

for index, center in enumerate(centers):

seqOfCenter = center[0] * numOfRows + center[1]

distance[index] = kernel[pixel, pixel] + kernel[seqOfCenter, seqOfCenter] - 2 * kernel[pixel, seqOfCenter]

# Pick the index of minimum distance as the cluster of the point

clusters[pixel] = np.argmin(distance)

return clusters
```

It is same to the init cluster of part 1.

3. Kernel K-means

```
# Kernel k-means.
currentClusters = clusters.copy()
count = 0
iteration = 100

while True:

# Compute new clusters.
numOfPixels = numOfRows * numOfCols
newClusters = kernelClustering(numOfPixels, numOfClusters, kernel, currentClusters)

# Get the image state.
imageState = getCurrentImageState(numOfRows, numOfCols, newClusters)

imageStates.append(imageState)

if np.linalg.norm((newClusters - currentClusters), ord=2) < 0.001 or count >= iteration:
break

currentClusters = newClusters.copy()
count += 1
```

It is same to the kernel k-means part of part 1.

4. Kernel Clustering

```
157 Odef kernelClustering(numOfPixels, numOfClusters, kernel, clusters):

# Get number of members in each cluster
numOfMembers = np.array([np.sum(np.where(clusters == c, 1, 0)) for c in range(numOfClusters)])

# Get sum of pairwise kernel distances of each cluster
pairwise = getSumOfPairwiseOistance(numOfPixels, numOfClusters, numOfMembers, kernel, clusters)

163 newClusters = np.zeros(numOfPixels, dtype=int)

165 for p in range(numOfPixels):
166 distance = np.zeros(numOfClusters)

167 for c in range(numOfClusters):
168 distance[c] += kernel[p, p] + pairwise[c]

179 # Get distance from given data point to others in the target cluster
171 distance[c] -= 2.0 / numOfMembers[c] * distToOthers
newClusters[p] = np.argmin(distance)

175 preturn newClusters
```

It is same to the kernel clustering of part 1.

- Part 3 (2,3-clusters / k-means++)
 - 1. Init Center

For the init center code of part 3, I choose the init center based on kmeans++ strategy. The function will return centers in array form.

2. Init Clusters

```
def initClusters(numOfRows, numOfCols, numOfClusters, kernel, initMode):

# Init centers.
centers = initCenters(numOfRows, numOfCols, numOfClusters, initMode)

# K-means.
numOfPixels = numOfRows * numOfCols
clusters = np.zeros(numOfPixels, dtype=int)

# for pixel in range(numOfPixels):
# Compute the distance of every pixel to all centers.
distance = np.zeros(numOfClusters)

# or index, center in enumerate(centers):
# seqOfCenter = center[0] * numOfRows + center[1]
# pick the index of minimum distance as the cluster of the point
clusters[pixel] = np.argmin(distance)

# return clusters
```

It is same to the init cluster of part 1.

3. Kernel K-means

It is same to the kernel k-means part of part 1.

4. Kernel Clustering

```
# Get number of members in each cluster

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numOfMembers = np.array([np.sum(np.where(clusters == c, 1, 0)) for c in range(numOfClusters)])

# Get sum of pairwise kernel distances of each cluster

pairwise = getSumOfPairwiseDistance(numOfPixels, numOfClusters, numOfMembers, kernel, clusters)

newClusters = np.zeros(numOfPixels, dtype=int)

for p in range(numOfFixels):

distance = np.zeros(numOfClusters)

for c in range(numOfClusters):

distance[c] += kernel[p, p] + pairwise[c]

# Get distance from given data point to others in the target cluster

distToOthers = np.sum(kernel[p, :][np.where(clusters == c)])

distance[c] -= 2.0 / numOffMembers[c] * distToOthers

newClusters[p] = np.argmin(distance)

return newClusters
```

It is same to the kernel clustering of part 1.

- Common Parts
 - 1. Output the GIF Results

```
# Output the gif result.

filename = f'./output/kennel_kmeans/kernel_kmeans_{index}_' \

filename = f'./output/kennel_kmeans/kernel_kmeans_{index}_' \

f'cluster_{numOfClusters}_' \

f''{"kmeans++" if initMode else "random"}.gif'

os.makedirs(os.path.dirname(filename), exist_ok=True)

imageStates[0].save(filename, save_all=True, append_images=imageStates[1:], optimize=False, loop=0, duration=100)
```

After converging, we will output the result with gif pics.

- 2. Spectral Clustering
 - Common Parts
 - 1. Load Data

The objective of the readImagesINpArray function is to load data from the input file, it will be called in the main function.

2. Compute Kernel

```
def computeKernel(image, gammaC):

# Get image shape.

rows, cols, colors = image.shape

# Compute the color distance.

numOfPixels = rows * cols

colorDist = cdist(image.reshape(numOfPixels, colors), image.reshape(numOfPixels, colors), 'sqeuclidean')

# Compute the indices of a grid.

indices = np.indices((rows, cols))

indicesOfRow = indices[0]

# Compute the indices vector.

indicesVector = np.hstack((indicesOfRow.reshape(-1, 1), indicesOfCol.reshape(-1, 1)))

# Compute the spatial distance.

spatialDist = cdist(indicesVector, indicesVector, 'sqeuclidean')

# The kernel formula in spec.

return np.multiply(np.exp(-gammaS * spatialDist), np.exp(-gammaC * colorDist))
```

We reuse the **computeKernel** function in kernel k-means program, so the explanation is same to the content in kernel k-means section.

- Part 1 (2-clusters / randomly init)
 - 1. Compute Matrix U

```
# Compute matrixU which containing eigenvectors.

# Compute degree matrixD, outHode, numOfclusters):

# Compute degree matrixD and Laplacian matrixL.

matrixD = np.zeros_like(matrixW):

matrixD[index, row in enumerate(matrixW):

matrixD[index, index] += np.sum(row)

matrixL = matrixD - matrixW

# Normalized cut.

# Compute the normalized Laplacian matrixL.

for idx in range(len(matrixD)):

matrixD[idx, idx] = 1.0 / np.sqrt(matrixD[idx, idx])

## Compute eigenvalues and eigenvectors.

## Compute eigenvectors = np.linalg.eig(matrixL)

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## Sort the eigenvalues and find indices of nonzero eigenvalues.

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```

According to the lecture PPT, we know that L = D - W. So we compute the <code>matrixD</code> first, and then compute the <code>matrixL</code> based on D and W. After obtaining the <code>matrixL</code>, we will based on the input parameter <code>cutMode</code> to determine if we need to normalize the <code>matrixL</code>. The we will find the eigenvectors (with to zero eigenvalues) as the return value (<code>matrixU</code>).

2. Spectral Clustering

We will use the matrixu to get the initial centers, and perform k-means to get the result cluster.

3. Init Center

```
69 def initCenters(numOfRows, numOfCols, numOfClusters, matrixU, initMode):
70 if initMode == 1:
71  # Random strategy.
72  numOfPixels = numOfRows * numOfCols
73 preturn matrixU[np.random.choice(numOfPixels, numOfClusters)]
```

For the init center code of part 1, I just initialize the center randomly according to the number of number of clusters and also the eigenspace.

4. K-means

```
# Kernel k-means.

currentCenters = centers.copy()

newClusters = np.zeros(numOfPixels, dtype=int)

count = 0

iteration = 100

while True:

# Compute new cluster.

newClusters = kmeansClustering(numOfPixels, numOfClusters, matrixU, currentCenters)

# Compute new centers.

newCenters = kmeansRecomputeCenters(numOfClusters, matrixU, newClusters)

# Get new state.

imageStates.append(getCurrentImageState(numOfRows, numOfCols, newClusters))

if np.linalg.norm((newCenters - currentCenters), ord=2) < 0.01 or count >= iteration:

break

# Update current parameters.

currentCenters = newCenters.copy()

count += 1
```

We perform k-means based on initial centers. In each round, we use kmeansClustering to get new cluster, we also use the kMeansRecomputeCenters to update the centers. We will calculate the difference between the currentCenters and newCenters to check if it is already converged.

5. Output the GIF Results

```
# Output the gif result.

filename = f'./output/spectral_clustering/spectral_clustering_{index}_' \

f'clusterinumOfcLusters}_' \

f'clusterinumOfcLusters}_-' \

f'*(means++" if initMode else "random"}_-' \

f'*(mormalized" if cutMode else "ratio"}.gif'

os.makedirs(os.path.dirname(filename), exist_ok=True)

if len(imageStates) > 1:

imageStates[0].save(filename, save_all=True, append_images=imageStates[1:], optimize=False, loop=0,

duration=100)

else:

imageStates[0].save(filename)
```

After converging, we will output the result with gif pics.

- Part 2 (3-clusters / randomly init)
 - 1. Compute Matrix U

It is same to the compute matrix U of part 1.

2. Spectral Clustering

It is same to the spectral clustering of part 1.

3. Init Center

```
69 def initCenters(numOfRows, numOfCols, numOfClusters, matrixU, initMode):
78 def initMode == 1:
71  # Random strategy.
72  numOfFixels = numOfRows * numOfCols
73 def return matrixU[np.random.choice(numOfFixels, numOfClusters)]
```

It is same to the init center of part 1.

4. K-means

It is same to the k-means of part 1.

5. Output the GIF Results

```
# Output the gif result.

filename = f'./output/spectral_clustering/spectral_clustering_{index}_' \
f'cluster/numOfClusters}_' \
f'c
```

It is same to the output results of part 1.

- Part 3 (2,3-clusters / k-means++)
 - 1. Compute Matrix U

```
# Compute matrixU which containing eigenvectors.

# Compute degree matrixU, cutHode, numOfCCLusters):

# Compute degree matrixD and Laplacian matrixL.

matrixD = np.zeros_like(matrixW)

for index, row in enumerate(matrixW):

matrixD[index, index] += np.sum(row)

matrixL = matrixD - matrixW

# Normalized cut.

# Compute the normalized Laplacian matrixL.

for idx in range(len(matrixD)):

matrixD[idx, idx] = 1.0 / np.sqrt(matrixD[idx, idx])

matrixL = matrixD.dot(matrixL).dot(matrixD)

# Compute eigenvalues and eigenvectors.

eigenvalues, eigenvectors = np.linalg.eig(matrixL)

# Sort the eigenvalues and find indices of nonzero eigenvalues.

sortedIdx = np.argsort(eigenvalues)

sortedIdx = sortedIdx[eigenvalues[sortedIdx] > 0]

return eigenvectors[sortedIdx[eigenvalues]].T
```

It is same to the compute matrix U of part 1.

2. Spectral Clustering

It is same to the spectral clustering of part 1.

3. Init Center

```
# K-means++ strategy.
# K-means++ strategy.
# Compute indices of a grid.
indices = np.indices((numOfRows, numOfCols))
indicesOfRow = indices[0]

# Compute the indices vector.
indicesVector = np.hstack((indicesOfRow.reshape(-1, 1), indicesOfCol.reshape(-1, 1)))

# Randomly pick first center.
numOfPixels = numOfRows * numOfCols
centers = [indices[np.random.choice(numOfPixels, 1)[0]].tolist()]

# Find remaining centers.
for _ in range(numOfCutsters - 1):
# Compute min distance for each point to all found centers.
distance = np.zeros(numOfPixels)

for index, indice in enumerate(indicesVector):
minDistance = np.Inf

for center in centers:
    dist = np.linds_nome(indice - center)
    minDistance = dist if dist < minDistance else minDistance
    distance[index] = minDistance

# Divide the distance by its sum to get probability.
distance /= np.sum(distance)
# Set a new center.
centers.append(indices[np.random.choice(numOfPixels, 1, p=distance)[0]].tolist())

# Change from index to feature index.
for index, center in enumerate(centers):
centers[index] = matrixU[center[0] * numOfRows + center[1], :]

return np.array(centers)
```

For the init center code of part 3, I choose the init center based on kmeans++ strategy. The function will return centers in array form based on the eigenspace (matrixu).

4. K-means

```
# Kernel k-means.
currentCenters = centers.copy()
newClusters = np.zeros(numOfPixels, dtype=int)
count = 0
iteration = 100

while True:
# Compute new cluster.
newClusters = kmeansClustering(numOfPixels, numOfClusters, matrixU, currentCenters)

# Compute new centers.
newCenters = kmeansRecomputeCenters(numOfClusters, matrixU, newClusters)

# Get new state.
imageStates.append(getCurrentImageState(numOfRows, numOfCols, newClusters))

if np.linalg.norm((newCenters - currentCenters), ord=2) < 8.01 or count >= iteration:
break

# Update current parameters.
currentCenters = newCenters.copy()
count += 1
```

It is same to the k-means of part 1.

5. Output the GIF Results

```
# Output the gif result.

filename = f'./output/spectral_clustering/spectral_clustering_{index}_' \

fictuster{numOfClusters}_' \

f'ctuster{numOfClusters}_' \

f't*means++" if initNode else "random"}_' \

f't*normalized" if cutNode else "ratio"}.gif'

os.makedirs(os.path.dirname(filename), exist_ok=True)

if len(imageStates) > 1:

imageStates[0].save(filename, save_all=True, append_images=imageStates[1:], optimize=False, loop=0,

duration=100)

else:

imageStates[0].save(filename)
```

It is same to the output results of part 1.

- Part 4 (examine points)
 - 1. Plot Result

```
def plotResult(matrixU, clusters, index, initMode, cutMode):
    colors = ['r', 'b']
    plt.clf()

for idx, point in enumerate(matrixU):
    pt.scatter(point[0], point[1], c=colors[clusters[idx]])

# Save the figure.

filename = f'./output/spectral_clustering/eigenspace_{index}_' \
    f'{"normalized" if cutMode else "ratio"}.png'
    os.makedirs(os.path.dirname(filename), oxist_ok=True)

plt.savefig(filename)
```

Capture the result to examine whether the data points within the same cluster do have the same coordinates in the eigenspace of graph Laplacian or not.

- Experiments & Discussion
 - 1. Kernel K-means
 - Image 1
 - 1. Result

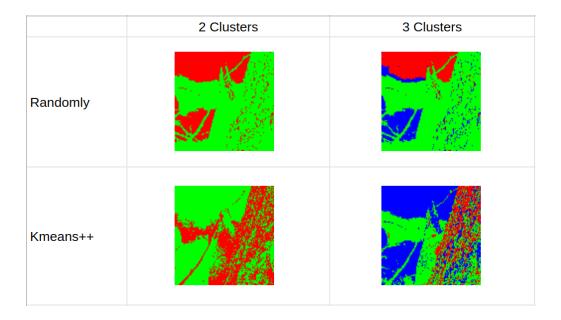
	2 Clusters	3 Clusters
Randomly		
Kmeans++		

2. Discussion

- Kmeans++ strategy can get better initial clustering.
- For image 1, the better value of k is 2 (sea and island).

• Image 2

1. Result

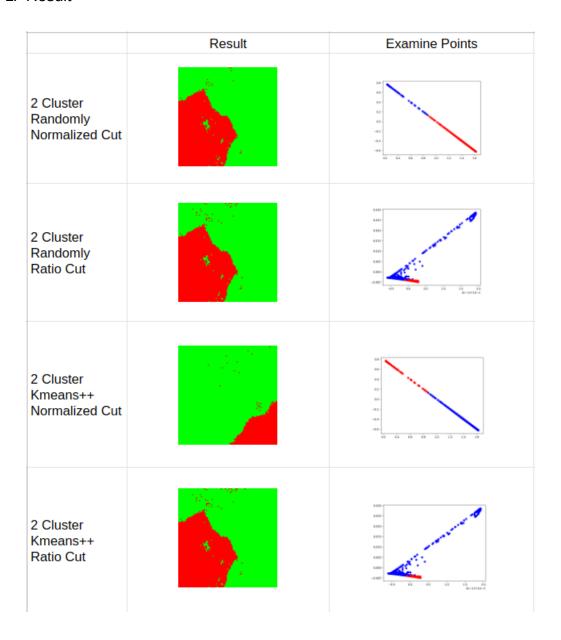


2. Discussion

• For image 2, the better value of k is 3 (tree, rabbit, and background).

2. Spectral Clustering

- Image 1 / 2 Clusters
 - 1. Result

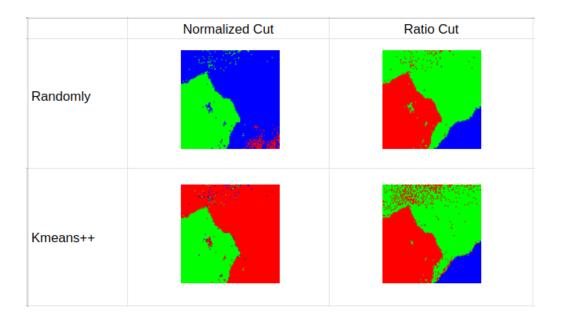


2. Discussion

• Kmeans++ strategy can get better initial clustering.

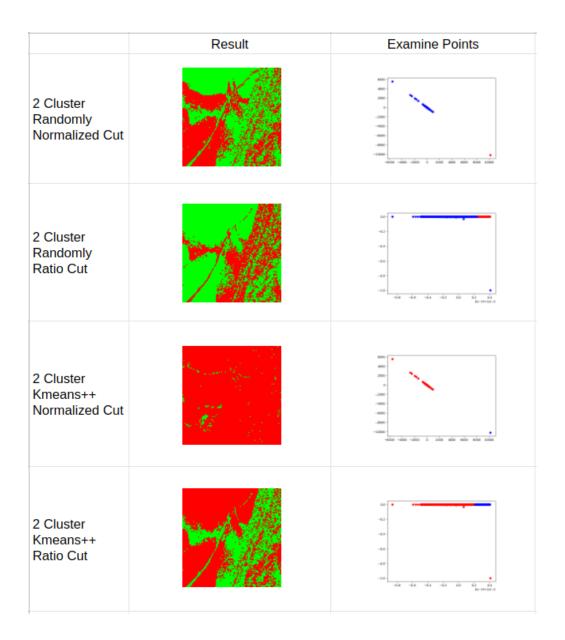
• Image 1 / 3 Clusters

1. Result



2. Discussion

- For image 1, the better value of k is 2 (sea and island). In randomly, normalized cut, we can see that there are only some of parts with red color.
- Image 2 / 2 Clusters
 - 1. Result



2. Discussion

- All of experiments result show that we can well separate the point in eigenspace.
- Image 2 / 3 Clusters
 - 1. Result

	Normalized Cut	Ratio Cut
Randomly		
Kmeans++		

2. Discussion

• The randomly init is not good initially (we can get the result in the randomly init, normalized cut).

Observations

- 1. How to choose k for supervised learning?
 - We may try the see the result first, and then decide the k value it should be. Because there is no general solution to find one.
- 2. Kmeans++ strategy is better than randomly initializing (the initial result).
- 3. We can see that it hardly classifies data points using normalized cut. Most of points are classified into same cluster.