Machine Learning Homework 6

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```
Environment

Code with detailed explanations
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Experiments settings and results & Discussion
   Part1
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   Part3
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Cobservations and discussion
```

Environment

Language: C++Standard: C++20

Code with detailed explanations

Libraries

Used library => Corresponding library in homework description

- Eigen => numpy
- OpenCV, matplotlib++ => visualization
- Boost => filesystem, sort
- OpenMP (for parallel acceleration)

Visualization

Main part

Use fittingHistory which records the labels for every iterations to make a video.

The duration of one iteration is 1s.

cv::addweighted function is used to combine the clustering results (segmentation mask) with original image.

drawMask function is used to convert the labels to segmentation mask.

```
kernelKMeans.fit(kernel);
const std::vector<Eigen::VectorXi> &fittingHistory = kernelKMeans.getFittingHistory();

writer.open(imageFile + "_video.mp4", codec, fps, image.size());
// check if we succeeded
if (!writer.isOpened())
{
    throw std::runtime_error("Could not open the output video file for write");
}

for (int i = 0; i < fps; i++)
{
    writer.write(image);
}

cv::Mat result;
cv::Mat mask;
for (std::size_t i = 1; i < fittingHistory.size(); i++)
{
    mask = drawMask(fittingHistory[i], numberOfClusters, image.cols, image.rows);
    cv::addWeighted(image, 0.5, mask, 0.5, 0, result);
    for (int i = 0; i < fps; i++)
    {
        writer.write(result);
    }
}

writer.release();

cv::imwrite(imageFile + "_mask.png", mask);
cv::imwrite(imageFile + "_final.png", result);
}

cv::imwrite(imageFile + "_final.png", result);</pre>
```

Draw mask

Convert the labels to segmentation mask.

Tabels contains the clustering results, from 0 to k-1.

cv::applyColorMap is used to map the labels to the corresponding color to avoid duplicate colors.

```
cv::Mat drawMask(const Eigen::VectorXi &labels, int numberOfClusters, unsigned int width, unsigned int height)
{
    Eigen::VectorX<unsigned char> maskData = labels.cast<unsigned char>();
    cv::Mat mask = cv::Mat(cv::Size(width, height), CV_8UC1, reinterpret_cast<void *>(maskData.data()));

mask *= (255 / numberOfClusters);

cv::Mat bgrMask;
    cv::cvtColor(mask, bgrMask, cv::COLOR_GRAY2BGR);
    cv::applyColorMap(bgrMask, bgrMask, cv::COLORMAP_COOL);
    return bgrMask;
}
```

Part1

- Kernel K-Means
 - Pseudo-code

Ignore weight

KERNEL_KMEANS_BATCH $(K, k, w, t_{max}, \{\pi_c^{(0)}\}_{c=1}^k, \{\pi_c\}_{c=1}^k)$

Input: K: kernel matrix, k: number of clusters, w: weights for each point, t_{max} : optional maximum number of iterations, $\{\pi_c^{(0)}\}_{c=1}^k$: optional initial clusters **Output:** $\{\pi_c\}_{c=1}^k$: final partitioning of the points

- 1. If no initial clustering is given, initialize the k clusters $\pi_1^{(0)}, ..., \pi_k^{(0)}$ (i.e., randomly). Set t = 0.
- 2. For each point \mathbf{a}_i and every cluster c, compute

$$d(\mathbf{a}_i, \mathbf{m}_c) = K_{ii} - \frac{2\sum_{\mathbf{a}_j \in \pi_c^{(t)}} w_j K_{ij}}{\sum_{\mathbf{a}_j \in \pi_c^{(t)}} w_j} + \frac{\sum_{\mathbf{a}_j, \mathbf{a}_l \in \pi_c^{(t)}} w_j w_l K_{jl}}{(\sum_{\mathbf{a}_j \in \pi_c^{(t)}} w_j)^2}.$$

3. Find $c^*(\mathbf{a}_i) = \operatorname{argmin}_c d(\mathbf{a}_i, \mathbf{m}_c)$, resolving ties arbitrarily. Compute the updated clusters as

$$\pi_c^{(t+1)} = \{ \mathbf{a} : c^*(\mathbf{a}_i) = c \}.$$

4. If not converged or $t_{max} > t$, set t = t + 1 and go to Step 2; Otherwise, stop and output final clusters

Main function

Arguments

path: the image data folder

numberOfClusters: the number of clusters

init: the selected initialization method, random or k-means++

gamma1, gamma2: the hyper-parameter of RBF kernel

```
oid run(const fs::path &path, int numberOfClusters, mlhw6::KMeansInitMethods init, double gamma1, double gar
       kernelKMeans.fit(kernel);
const std::vector<Eigen::VectorXi> &fittingHistory = kernelKMeans.getFittingHistory();
```

kmeans.h header: k-means related classes

Follow the scikit-learn logic design.

```
enum KMeansInitMethods

Random,
Random,
Reansplusplus,
};

class BasekMeans

public;

BasekMeans(int numberOfclusters, kMeansInitMethods init - KMeansInitMethods::Kmeansplusplus, int maximumEpochs - 200, int seed - 1234);

virtual tigen::WectorXi predict(const Eigen::MatrixXio 6x) - 0;

virtual tigen::WectorXi predict(const Eigen::Mefrconst Eigen::MatrixXio 6x);

const std::vector<Eigen::WectorXi pedict(const Eigen::Refcconst Eigen::MatrixXio 6x);

protected:
Eigen::WectorXi initializeCenters(const Eigen::Refcconst Eigen::MatrixXio 6x, KMeansInitMethods init, int seed, bool precomputed - false) const;

int maximumEpochs;
int maximumEpochs;
int maximumEpochs;
int maximumEpochs;
int maximumEpochs
int maximu
```

KernelkMeans constructor

```
KernelKMeans::KernelKMeans(int numberOfClusters, int maximumEpochs, int seed, KMeansInitMethods init): BaseKMeans(numberOfClusters, maximumEpochs, seed, init)

| Respective for the seed of the seed
```

preprocess function: extract RGB values and coordinates

Arguments

image: (H, W, 3), BGR values

o calculateKernel function: calculate all kernel values

```
Arguments pixels: (N, 3), RGB values coordinates: (N, 2), coordinates gamma1: \gamma_c scalar gamma2: \gamma_s scalar Formula
```

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

- S(x) is the spatial information (coordinate)
- C(x) is the color information (RGB)

```
Si verial Eigen::MatrixXd calculateKernel(const Eigen::MatrixX3d &pixels, const Eigen::MatrixX2i &coordinates, double gamma1, double gamma2)

auto colorKernel = mlhw6::rbf(pixels, pixels, gamma1);

const Eigen::MatrixX2d &fmp = coordinates.cast<double>();

auto coordinateKernel = mlhw6::rbf(tmp, tmp, gamma2);

return colorKernel.cwiseProduct(coordinateKernel);

return colorKernel.cwiseProduct(coordinateKernel);
```

o rbf function: RBF kernel

```
Arguments x_1: x vector x_2: x' vector y_3: x' vector y_4: x' vector y_5: x' vector y_6: x' vec
```

kernelkMeans.fit function: fit the data

Previously, we already calculated the kernel.

So, we use the precomputed kernel values directly.

Arguments

x: (N, N), the kernel values (gram matrix, similarity matrix)

Steps

- 1. Pick k centers
- 2. Calculate the cost between the data and centers
- 3. Assign the label which has the smallest distance to the data
- 4. Keep repeating 2, 3 step until the labels are not changed

fittingHistory is used to store the labels for every iterations.

Note: At the line 183, we do 1-x to get the distance matrix.

```
void KernelKMeans::fit(const Eigen::Ref<const Eigen::MatrixXd> 6x)
{
this→fittingHistory = std::vector<Eigen::VectorXi>{Eigen::VectorXi::Constant(x.rows(), -1)};

// x is similarity matrix (gram matrix)
auto centers = this→initializeCenters(1 - x.array(), this→init, this→seed, true);
this→fittingHistory.back()(centers).setLinSpaced(0, this→numberOfClusters - 1);

int epoch = 0;
bool sameLabels = false;
do
{
this→fittingHistory.push_back(this→assignLabels(x));

sameLabels = (this→fittingHistory[epoch].array() = this→fittingHistory[epoch + 1].array()).all();
epoch+;
} while (!sameLabels && epoch < this→maximumEpochs);

std::cout < "Finished at " < epoch < " epoch" < std::endl;
}

**This int epoch = 0;
bool sameLabels = false;
do
{
this→fittingHistory.push_back(this→assignLabels(x));
sameLabels = (this→fittingHistory[epoch].array() = this→fittingHistory[epoch + 1].array()).all();
epoch+;
} while (!sameLabels && epoch < this→maximumEpochs);

std::cout < "Finished at " < epoch < " epoch" < std::endl;
```

assignLabels function: calculate the cost and assign labels

Arguments

x: (N, N), the kernel values

Steps

1. Calculate the cost between the data and centers

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

2. Assign the label which has the smallest distance to the data

```
Eigen::VectorXi KernelKMeans::assignLabels(const Eigen::Ref<const Eigen::MatrixXd> &x) const

{

Eigen::MatrixXd distance(x.rows(), this→numberOfClusters);

momp parallel for
for (int k = 0; k < this→numberOfClusters; k++)

{

Eigen::VectorXd selector = this→fittingHistory.back().cwiseEqual(k).cast<double>();
auto numberOfXInKCluster = selector.sum();
Eigen::MatrixXd xToKCluster = x * selector.asDiagonal();

Eigen::VectorXd secondTerm = 2 * xToKCluster.rowwise().sum() / numberOfXInKCluster;
auto thirdTerm = (selector.asDiagonal() * xToKCluster).sum() / std::pow(numberOfXInKCluster, 2);
distance(Eigen::all, k) = (x.diagonal() - secondTerm).array() + thirdTerm;
}

std::vector<int> labels(x.rows());

pragma omp parallel for
for (Eigen::Index i = 0; i < x.rows(); i++)
{
    // find nearest neighbor
    distance.row(i).minCoeff(&labels[i]);
}
return Eigen::VectorXi::Map(labels.data(), labels.size());
}
```

Spectral Clustering

- o Pseudo-code
 - Spectral Clustering (ratio cut)

```
Unnormalized spectral clustering
```

Input: Similarity matrix $S \in \mathbb{R}^{n \times n}$, number k of clusters to construct.

- \bullet Construct a similarity graph by one of the ways described in Section 2. Let W be its weighted adjacency matrix.
- ullet Compute the unnormalized Laplacian L.
- Compute the first k eigenvectors u_1, \ldots, u_k of L.
- ullet Let $U\in\mathbb{R}^{n imes k}$ be the matrix containing the vectors u_1,\dots,u_k as columns.
- ullet For $i=1,\ldots,n$, let $y_i\in\mathbb{R}^k$ be the vector corresponding to the i-th row of U.
- \bullet Cluster the points $(y_i)_{i=1,\dots,n}$ in \mathbb{R}^k with the k-means algorithm into clusters C_1,\dots,C_k .

Output: Clusters A_1,\ldots,A_k with $A_i=\{j|\ y_j\in C_i\}$.

Normalized Spectral Clustering (normalized cut)

Normalized spectral clustering according to Shi and Malik (2000)

Input: Similarity matrix $S \in \mathbb{R}^{n \times n}$, number k of clusters to construct.

- \bullet Construct a similarity graph by one of the ways described in Section 2. Let W be its weighted adjacency matrix.
- ullet Compute the unnormalized Laplacian L.
- Compute the first k generalized eigenvectors u_1, \ldots, u_k of the generalized eigenproblem $Lu = \lambda Du$.
- Let $U \in \mathbb{R}^{n \times k}$ be the matrix containing the vectors u_1, \dots, u_k as columns.
- ullet For $i=1,\ldots,n$, let $y_i\in\mathbb{R}^k$ be the vector corresponding to the $i ext{-th}$ row of U.
- \bullet Cluster the points $(y_i)_{i=1,\dots,n}$ in \mathbb{R}^k with the k-means algorithm into clusters C_1,\dots,C_k .

Output: Clusters A_1, \ldots, A_k with $A_i = \{j | y_j \in C_i\}$.

Main function

Arguments path: the image data folder model: Spectral Clustering instance numberOfClusters: the number of clusters gamma1, gamma2: the hyper-parameter of RBF kernel

```
switch (type)
{
   case SpectralClusteringType::Unnormalized:
   {
     auto model = mlhw6::SpectralClustering(numberOfClusters, init);
     run(path, model, numberOfClusters, gamma1, gamma2);
     break;
}
case SpectralClusteringType::Normalized:
   {
     auto model = mlhw6::NormalizedSpectralClustering(numberOfClusters, init);
     run(path, model, numberOfClusters, gamma1, gamma2);
     break;
}
default:
   std::cerr << "Unknown spectral clustering type." << std::endl;
   return 1;
}</pre>
```

```
void run(const fs::path &path, mlhw6::BaseSpectralClustering &model, int numberOfClusters, double gamma1, double gamma2)
{
    int fps = 30;
    int codec = cv::VideoWriter::fourcc('m', 'p', '4', 'v');
    cv::VideoWriter writer;

    for (auto imageFile : IMAGE_FILES)

    std::cout « imageFile « std::endl;

    auto fileName = (path / imageFile).generic_string();

    // read image
    auto image = cv::imread(fileName, cv::ImreadModes::IMREAD_COLOR);

    // extract RGB values and coordinates
    auto [pixels, coordinates] = preprocess(image);

    // calculate kernel
    auto kernel = calculateKernel(pixels, coordinates, gamma1, gamma2);

    model.fit(kernel);
    const std::vector<Eigen::VectorXi> &fittingHistory = model.getFittingHistory();

    writer.open(imageFile + " video.mp4", codec, fps. image.size());
}
```

o spectral.h header: spectral clustering related classes

Follow the scikit-learn logic design.

```
class BaseSpectralClustering {
    public:
        BaseSpectralClustering(int numberOfClusters, KMeansInitMethods init = KMeansInitMethods::Kmeansplusplus, int maximumEpochs = 200, int seed = 1234);

    virtual void fit(const Eigen::Ref<const Eigen::MatrixXd> 6x) = 0;

    virtual Eigen::VectorXi fitAndBredict(const Eigen::Ref<const Eigen::MatrixXd> 6x);

    const std::VectorCigen::VectorXi> 6getFittingHistory() const;

    const Eigen::MatrixXd bgetEigenMatrix() const;

    protected:
        KMeans KMeans;
        Eigen::MatrixXd eigenMatrix;

        int numberOfClusters;
        unsigned int numberOfThreads;
    };

    class SpectralClustering::BaseSpectralClustering;
    void fit(const Eigen::Ref<const Eigen::MatrixXd> 6x) override;
};

class NormalizedSpectralClustering::BaseSpectralClustering;
    void fit(const Eigen::Ref<const Eigen::MatrixXd> 6x) override;
};

void fit(const Eigen::Ref<const Eigen::MatrixXd> 6x) override;
};

void fit(const Eigen::Ref<const Eigen::MatrixXd> 6x) override;
};
```

kmeans.h header: k-means related classes

Follow the scikit-learn logic design.

```
cmum KMeansInitMethods

Random,
Random
```

o preprocess function: extract RGB values and coordinates

Arguments

image: (H, W, 3), BGR values

o calculateKernel function: calculate all kernel values

```
Arguments pixels: (N, 3), RGB values coordinates: (N, 2), coordinates gamma1: \gamma_c scalar gamma2: \gamma_s scalar Formula k(x,x') = e^{-\gamma_s \|S(x) - S(x')\|^2} \times e^{-\gamma_c \|C(x) - C(x')\|^2}
```

S(x) is the spatial information (coordinate)

C(x) is the color information (RGB)

o rbf function: RBF kernel

```
Arguments  \hbox{x1: } x \hbox{ vector} \\ \hbox{x2: } x' \hbox{ vector} \\ \hbox{gamma: } \gamma \hbox{ scalar} \\ \hbox{Formula} \\ k(x,x')=e^{-\gamma||x-x'||^2}
```

```
template <typename DerivedA, typename DerivedB, typename Out = Eigen::Matrix<double, DerivedA::RowsAtCompileTime, DerivedB::RowsAtCompileTime>> Out rbf(const Eigen::MatrixBase<DerivedA> 5x1, const Eigen::MatrixBase<DerivedB> 6x2, double gamma)

{
    Out result(x1.rows(), x2.rows());
    #pragma omp parallel for
    for (Eigen::Index i = 0; i < x1.rows(); i++)
    {
        result.row(i) = (-gamma * (x2.rowwise() - x1.row(i)).rowwise().squaredNorm().transpose()).array().exp();
    }
    return result;
}
```

model.fit function: fit the data

Previously, we already calculated the kernel.

So, we use the precomputed kernel values directly.

Arguments

x: (N, N), the kernel values (gram matrix, similarity matrix)

■ Spectral Clustering (ratio cut)

Algorithm

Unnormalized spectral clustering

Input: Similarity matrix $S \in \mathbb{R}^{n \times n}$, number k of clusters to construct.

- \bullet Construct a similarity graph by one of the ways described in Section 2. Let W be its weighted adjacency matrix.
- ullet Compute the unnormalized Laplacian L.
- Compute the first k eigenvectors u_1, \ldots, u_k of L.
- ullet Let $U\in\mathbb{R}^{n imes k}$ be the matrix containing the vectors u_1,\dots,u_k as columns.
- ullet For $i=1,\ldots,n$, let $y_i\in\mathbb{R}^k$ be the vector corresponding to the i-th row of U.
- ullet Cluster the points $(y_i)_{i=1,\dots,n}$ in \mathbb{R}^k with the k-means algorithm into clusters C_1,\dots,C_k .

Output: Clusters A_1,\ldots,A_k with $A_i=\{j|\ y_j\in C_i\}$.

Normalized Spectral Clustering (normalized cut)

Algorithm

```
Normalized spectral clustering according to Shi and Malik (2000) Input: Similarity matrix S \in \mathbb{R}^{n \times n}, number k of clusters to construct.

• Construct a similarity graph by one of the ways described in Section 2. Let W be its weighted adjacency matrix.

• Compute the unnormalized Laplacian L.

• Compute the first k generalized eigenvectors u_1, \ldots, u_k of the generalized eigenproblem Lu = \lambda Du.

• Let U \in \mathbb{R}^{n \times k} be the matrix containing the vectors u_1, \ldots, u_k as columns.

• For i = 1, \ldots, n, let y_i \in \mathbb{R}^k be the vector corresponding to the i-th row of U.

• Cluster the points (y_i)_{i=1,\ldots,n} in \mathbb{R}^k with the k-means algorithm into clusters C_1, \ldots, C_k.

Output: Clusters A_1, \ldots, A_k with A_i = \{j \mid y_j \in C_i\}.
```

kMeans.fit function: fit the data

Use the points in eigen space to perform k-means clustering.

Steps

1. Pick k centers

- 2. E step
 - 1. Calculate the cost between the data and centers
 - 2. Assign the label which has the smallest distance to the data
- 3. M step

Recalculate the centers by averaging the points which belong to the same cluster.

4. Keep repeating 2, 3 step until the labels are not changed

o assignLabels function: calculate the cost and assign labels

Arguments
x: (N, N), the kernel values
Steps

- 1. Calculate the cost (Euclidean distance) between the data and centers
- 2. Assign the label which has the smallest distance to the data

Part2

Use command to control the number of clusters and other parameters.

Kernel K-Means

init0 => use random initialization1 => use k-means++ initialization

```
int main(int argc, char *argv[])

if (argc < 6)
{
    std::cerr << "Usage: " << argv[0] << " <data path> <number of cluster> <init> <gamma1> <gamma2>" << std::endl;
    return 1;
}</pre>
```

• Spectral Clustering

```
spectral clustering type
0 => use spectral clustering algorithm
1 => use normalized clustering algorithm
init
0 => use random initialization
1 => use k-means++ initialization
```

Part3

• initializeCenters function: pick k centers initialized by the selected method

```
Arguments
```

x: (N, N) precomputed distance matrix or (N, features) data init: the selected initialization method, random or k-means++ seed: the random seed precomputed: x is the precomputed distance matrix or not.

• Random initialization

```
Arguments
```

x: (N, N) precomputed distance matrix or (N, features) data numberOfClusters: k clusters seed: the random seed

Steps

- 1. generate the sequence of indexes, 0 ~ N-1
- 2. shuffle the sequence
- 3. pick the top k rows as the centers

```
Eigen::VectorXi randomInitialization(const Eigen::Ref<const Eigen::MatrixXd> &x, int numberOfClusters, int seed)

auto rng = std::mt19937_64(seed);
std::vector<int> sequence(x.rows());
std::iota(sequence.begin(), sequence.end(), 0);
std::shuffle(sequence.begin(), sequence.end(), rng);

Eigen::Map<Eigen::VectorXi> tmp = Eigen::VectorXi::Map(sequence.data(), sequence.size());
return tmp.topRows(numberOfClusters);

}
```

• K-Means++ initialization

Arguments

x: (N, N) precomputed distance matrix or (N, features) data numberOfClusters: k clusters seed: the random seed

Steps

- 1. Choose one center uniformly at random among the data points.
- 2. For each data point x not chosen yet, compute $D(x)^2$ (the squared Euclidean distance) or use the precomputed distance matrix, the distance between x and the nearest center that has already been chosen.
- 3. Choose one new data point at random as a new center, using a weighted probability distribution where a point x is chosen with probability proportional to $D(x)^2$. (The farthest point will be chosen.)

```
Eigen::WetOrXi MeansPlusPlusInitialization(const Eigen::Mefrconst Eigen::MatrixXd> 8x, int numberOfClusters, int seed, bool precomputed)
{
    auto rmg - std::m19907_66(seed);
    std::m19907_66(seed);
    std::m19907_66(seed);
    // 1. Choses one center uniformly at random among the data points.
    // 2. (red cain-ren') (0, rms = 1);
    auto sampler * std::m16rm_int,distribution(0, static_castcint>(x.tors()) - 1);
    candidates.push_beck(sampler(rmg));
    mnberOfClusters -:
    // 2. (red cain-ren');
    // 3. (red cain-ren');
    // 4. (red cain-ren');
    // 5. (red cain-ren');
    // 6. (red cain-ren');
    // 6. (red cain-ren');
    // 7. (red cain-ren');
    // 8. (red cain-ren');
    // 8. (red cain-ren');
    // 8. (red cain-ren');
    // 7. (red cain-ren');
    // 8. (red cain-ren');
    // 9. (red ca
```

Part4

Visualize the eigen space.

```
void plotEigenSpace(const std::string &path, const Eigen::MatrixXd &eigenMatrix)
{
    matplot::figure();

const Eigen::VectorXd &vecX = eigenMatrix.col(0);
    const Eigen::VectorXd &vecY = eigenMatrix.col(1);

std::vector<double> x(vecX.begin(), vecX.end());
    std::vector<double> y(vecY.begin(), vecY.end());
    matplot::scatter(x, y);
    matplot::save(path);
}
```

Experiments settings and results & Discussion

Part1

• Kernel K-Means

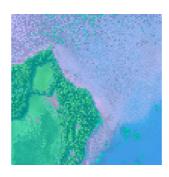
Number of clusters: 2

Initialization method: random

Gamma1: 0.00001 Gamma2: 0.00001

o image1

- <u>Video</u>
- Final

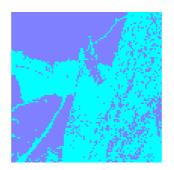


Mask



- o image2
 - <u>Video</u>
 - Final





• Spectral Clustering

Part2

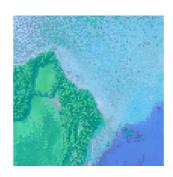
• Kernel K-Means

Initialization method: random

Gamma1: 0.00001

Gamma2: 0.00001

- o K = 3 ⋅ image1
 - <u>Video</u>
 - Final



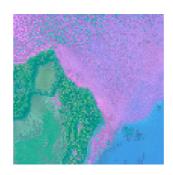


- o K = 3 ⋅ image2
 - <u>Video</u>
 - Final

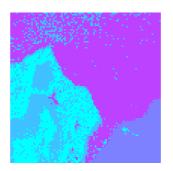




- o K = 4 · image1
 - <u>Video</u>
 - Final

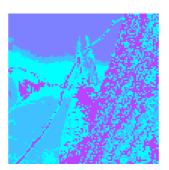


Mask



- o K = 4 · image2
 - <u>Video</u>
 - Final





Part3

• Kernel K-Means

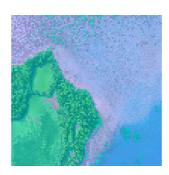
I think there is no significant difference between random and k-means++.

Initialization method: k-means++

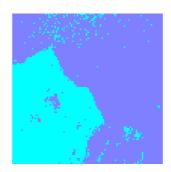
Gamma1: 0.00001

Gamma2: 0.00001

- o K = 2 ⋅ image1
 - <u>Video</u>
 - Final



Mask



- o K = 2 ⋅ image2
 - <u>Video</u>
 - Final





- ∘ K = 3 · image1
 - <u>Video</u>
 - Final



Mask



- o K = 3 ⋅ image2
 - <u>Video</u>
 - Final

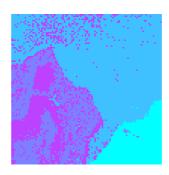




- o K = 4 · image1
 - <u>Video</u>
 - Final



Mask



- \circ K = 4 · image2
 - <u>Video</u>
 - Final



Mask



• Spectral Clustering

Part4

Observations and discussion

• Coming soon...