SPIRAL Dataset k=10

```
clear
clc
```

Load the file

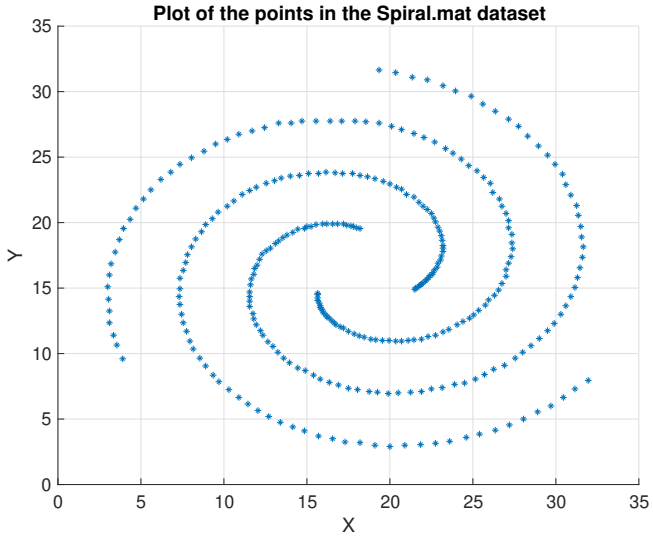
```
spiral_mat = load('Spiral.mat');

% Display the structure of the file
disp(spiral_mat);
```

```
    X: [312x3 double]
```

```
% Extract the matrix of points
% X_tot contains the right labels of the clusters as third column
X_tot = spiral_mat.X;
X = X_tot(:, 1:2);
```

Plot the points

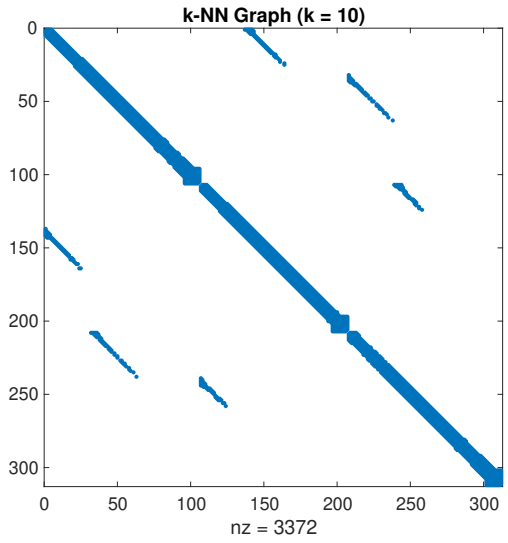
```
figure;
scatter(X(:,1), X(:,2), 10, Marker='*');
xlabel('X');
ylabel('Y');
title('Plot of the points in the Spiral.mat dataset');
grid on;
```



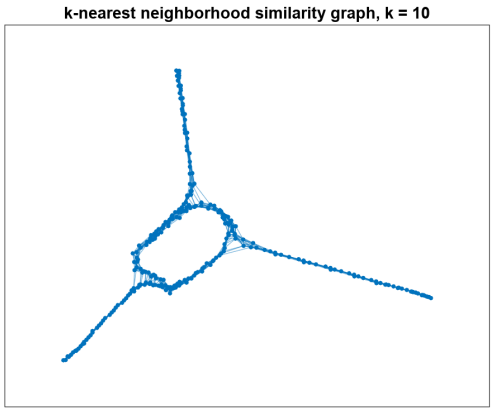
1. Similarity matrix and adjacency matrix

Three independent simulations of the script were carried out, with k set to 10, 20 and 40, respectively. The results presented in this script were obtained with a value of k fixed at 10. To analyse other values of k, it is simply necessary to change the value of k in the source code and re-run the script.

```
k_values = [10, 20, 40];
k = 10;

% Construct the k-nearest neighborhood similarity graph and its adjacency
% matrix W
W = knn_graph(X, k);

% Visualize the graph using its similarity matrix
figure;
spy(W);
title(['k-NN Graph (k = ', num2str(k), ')']);
```




```
% Store W as a sparse matrix
W = sparse(W);

% Visualize the graph G corresponding to the adjacency matrix W
G = graph(W);
figure;
plot(G);
title(['k-nearest neighborhood similarity graph, k = ', num2str(k)]);
```

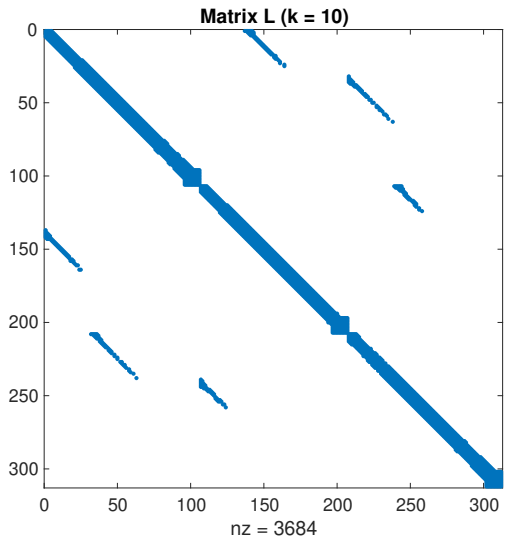


2.Construct the degree matrix D and the Laplacian matrix L

```
N = size(W, 1);

% Inizialize the degree matrix D
D = zeros(N, N);

% The degree of each point is given by the sum of the elements of each row in W
D = diag(sum(W, 2));

% Save D in a sparse format
D = sparse(D);

% Compute the Laplacian matrix L
L = D - W;

if issparse(L) % issparse(L)==1 means that L is stored in a sparse format
    disp("The matrix L is stored in a sparse format")
end
```

```
The matrix L is stored in a sparse format
```

```
% Plot the Laplacian matrix L
figure;
spy(L);
title(['Matrix L (k = ', num2str(k), ')']);
```



3.Compute the number of connected components of the similarity graph

```
% The points with the same number belong to the same connected component
bins = conncomp(G);

% Number of connected components
num_components = max(bins);

% Display the result
disp(['Number of connected components: ', num2str(num_components)]);
```

```
Number of connected components: 1
```

4 - 5. Compute eigenvalues and eigenvectors

This script implements two methods for computing the M smallest eigenvalues and their corresponding eigenvectors: the deflation method and the inverse power method. Currently, the script is configured to use the deflation method. The first eigenvector is manually set as $eigvects(:,1) = \frac{ones(N,1)}{norm(ones(N,1))}$, reflecting the known eigenvector of all ones for the Laplacian matrix. Alternatively, the first eigenvalue and eigenvector can be computed using the inverse power method by uncommenting the line $[eigvalues(1), eigvects(:,1)] = invpower\_method(L, v(1:end), maxIter, relTol)$ and commenting out the line that manually sets the first eigenvector.

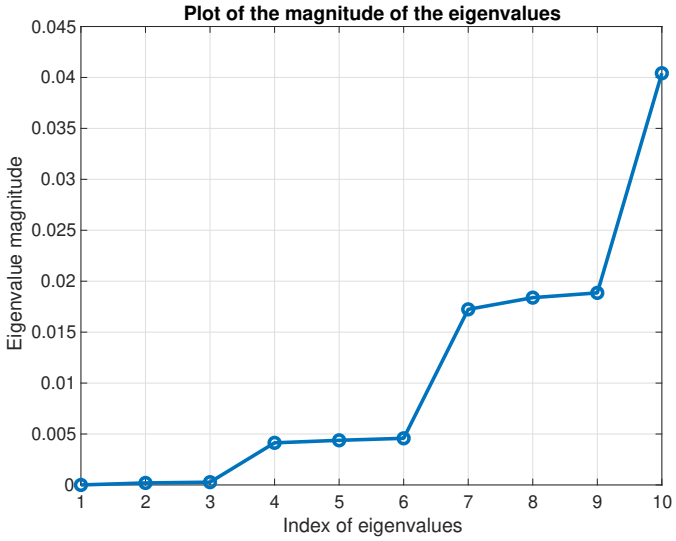
```
% Set a number M of values to be computed (later it will be changed)
M = 10;

% Inizialize the eigenvalues vector and the eigenvectors matrix
eigvalues = zeros(M, 1);
eigvects = zeros(N, M);

% Choose the vector v that will be used for the inverse power method
v = 0.5 * ones(N, 1);
v(1:2:N) = -0.5;
```

```
% Max iterations in the power method
maxIter = 1000;
% Relative tolerance
relTol = 1e-10;

% A known fact from theory is that L is semi pos def and has at least one
% eigenvalue = 0 and that the vector of all ones is a corresponding
% eigenvector
% eigvalues(1) = 0;
eigvects(:, 1) = ones(N,1)/ norm(ones(N,1));

% Or use the inverse power method to compute them
%[eigvalues(1), eigvects(:, 1)] = invpower_method(L, v(1:end), maxIter, relTol);

% Compute the reamining eigenvetors and eigenvalues
[eigvalues, eigvects, residualnorms] =
    deflation_method(L, v, eigvects, eigvalues,M, maxIter, relTol);

% Check how good the approximation is by comparing with eigs function of
% Matlab
[mat_eigvects, mat_eigs] = eigs(L, M, 'smallestabs');
norm(eigvalues - diag(mat_eigs))
```

```
ans = 1.2790e-10
```

Now, find the actual number M of eigenvalues that will be used for the clustering algorithm

```
% Plot the computed eigenvalues
x = 1:M;

figure
plot(x, eigvalues, '-o', 'LineWidth', 2);
xlabel('Index of eigenvalues');
ylabel('Eigenvalue magnitude');
title('Plot of the magnitude of the eigenvalues');
grid on;
```



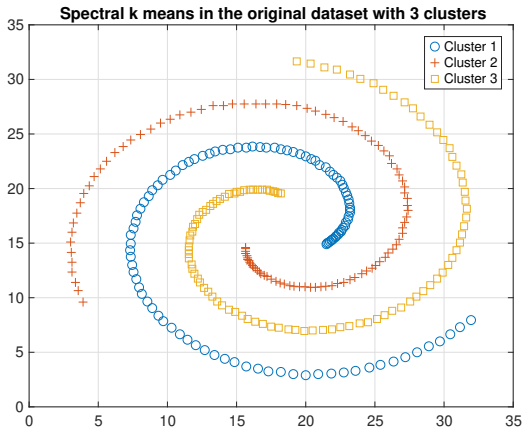
The suitable number of eigenvalues is 3 since eig4 is much larger than eig3 for k=10

```
M = 3; % K 40, maybe also 4

% Define the matrix U that will be used for the spectral clustering
U = eigvects(:, 1:M);
```

6 - 7 - 8. Spectral clustering, k means

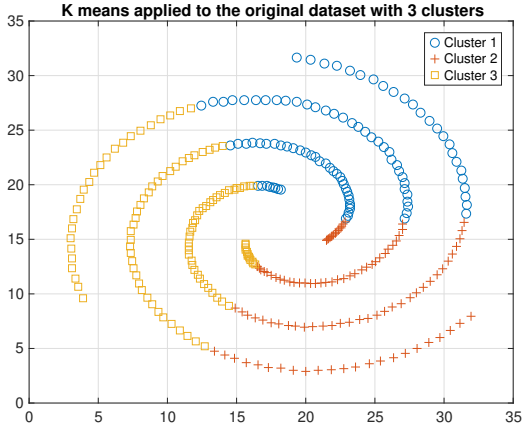
```
% Clusterize using k means and obtain the indices (and the centroids)
% inside the clusters of each point
[idx, C] = kmeans(U, M);

% Assing the original data to the corresponding clusters
A = cell(M, 1);

for i = 1:N
    % Find the cluster of y_i
    cluster_idx = idx(i);

    % Assing it to x_i
    A{cluster_idx} = [A{cluster_idx}; X(i, :)];
end

% Plot of the clusterized data in the original space
X_spect_clust = zeros(N, 3);
X_spect_clust(:, 1: 2) = X;
X_spect_clust(:, 3) = idx;

figure;
markers = ['o', '+', 's'];
gscatter(X_spect_clust(:,1), X_spect_clust(:,2), X_spect_clust(:, 3), [], markers, [], 5);
```

```
title(['Spectral k means in the original dataset with ', num2str(M), ' clusters']);
legend('Cluster 1', 'Cluster 2', 'Cluster 3');
grid on;
```



9.a K MEANS TO THE ORIGINAL DATA

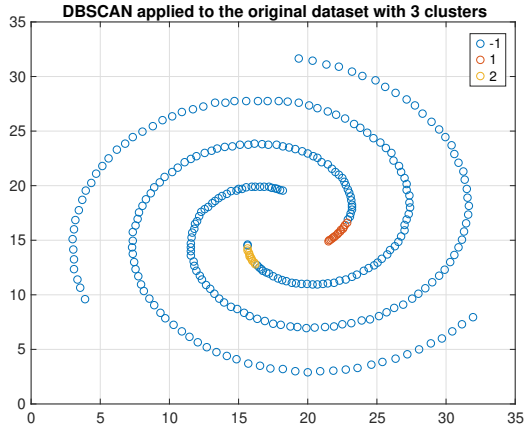
```
k_value = 3;  % 3

% Clusterize the original data
[idx_k, C_k] = kmeans(X, k_value);

% Add the index to X_kmeans
X_kmeans = zeros(N, 3);
X_kmeans(:, 1: 2) = X;
X_kmeans(:, 3) = idx_k;

figure;
markers = ['o', '+', 's'];
gscatter(X_kmeans(:,1), X_kmeans(:,2), X_kmeans(:, 3), [], markers, [], 5);
title(['K means applied to the original dataset with ', num2str(k_value), ' clusters']);
legend('Cluster 1', 'Cluster 2', 'Cluster 3');
grid on;
```



9.b DBSCAN TO THE ORIGINAL DATA

Apply the DBSCAN algorithm

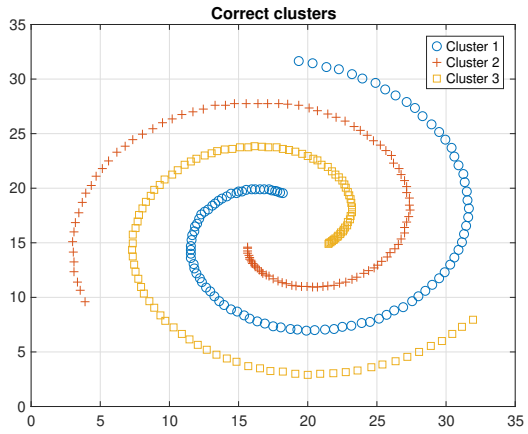
```
% Neighborhood radius
epsilon = 0.5;
% Minimum points for a cluster
minPts = 6;

idx_d = dbscan(X, epsilon, minPts);

X_dbscan = zeros(N, 3);
X_dbscan(:, 1: 2) = X;
X_dbscan(:, 3) = idx_d;

figure;
gscatter(X_dbscan(:,1), X_dbscan(:,2), X_dbscan(:, 3), [], 'o', 5);
title(['DBSCAN applied to the original dataset with ', num2str(M), ' clusters']);
grid on;
```



Difference with original clusters

Plot the original clusters

```
figure;
markers = ['o', '+', 's'];
gscatter(X_tot(:,1), X_tot(:,2), X_tot(:, 3), [], markers, [], 5);
title('Correct clusters');
legend('Cluster 1', 'Cluster 2', 'Cluster 3');
grid on;
```




```
% Compute the difference with the ones given by spectral clustering

% Shift the numbers in X_spect_clust(:,3) to have the same names as in
% X_tot

first_value = X_spect_clust(1, 3);
second_value = X_spect_clust(160, 3);
third_value = X_spect_clust(310, 3);

for i = 1:length(X_spect_clust(:, 3))
    if X_spect_clust(i, 3) == first_value
        X_spect_clust(i, 3) = 3;
    elseif X_spect_clust(i, 3) == second_value
        X_spect_clust(i, 3) = 1;
    elseif X_spect_clust(i, 3) == third_value
        X_spect_clust(i, 3) = 2;
    end
end

% Compute the difference
diff_clust = sum((X_tot(:,3) - X_spect_clust(:,3)))
```

```
diff_clust = 0
```