Exercise 2

1. Density Based Clustering: DENCLUE Write a script to implement the DENCLUE density-based clustering algorithm Algorithm 15.2 in chapter 15. The script should take as input a dataset D, the minimum density ξ, the tolerance for convergence ε, and the width h. Do not make any assumptions about the data (i.e., column names, etc), except that the last column gives the "true" cluster id.

Run your script on the iris.txt dataset, with ϵ =0.0001. Your script should output the following:

The number of clusters, and the size of each cluster. The density attractor, followed by the set of point in that cluster. Purity of the clustering, based on the true id. For Iris, you should use a value of ξ that gives you 3 clusters in the end, i.e., try different values and then finally report only the results for the value that gives you 3 clusters, since there are 3 true clusters in the data. Select the value of h empirically.

To speed up the computation for estimating the density at a point, you may want to first identify the K nearest neighbors, and use only those neighbors.

```
In [29]: import pandas as pd
    import numpy as np
    import networkx as nx
    from scipy.spatial import distance
    from sklearn.base import BaseEstimator, ClusterMixin
    print("Import Complete")

Import Complete
In [30]: FILE_NAME = "iris.txt"
```

```
In [31]: | ## Code adapted from @author: mgarrett
         def _hill_climb(x_t, X, W=None, h=0.3, eps=0.0001):
             This function climbs the 'hill' of the kernel density function
             and finds the 'peak', which represents the density attractor
             error = 99.
             prob = 0.
             x 11 = np.copy(x t)
             #Sum of the last three steps is used to establish radius
             #of neighborhood around attractor. Authors suggested two
             #steps works well, but I found three is more robust to
             #noisey datasets.
             radius_new = 0.
             radius old = 0.
             radius_twiceold = 0.
             iters = 0.
             while True:
                 radius_thriceold = radius_twiceold
                 radius twiceold = radius old
                 radius old = radius new
                 x_10 = np.copy(x_11)
                 x_11, density = _step(x_10, X, W=W, h=h)
                 error = density - prob
                 prob = density
                 radius new = np.linalg.norm(x 11-x 10)
                 radius = radius thriceold + radius twiceold + radius old + radius new
                 iters += 1
                 if iters>3 and error < eps:</pre>
                     break
             return [x_l1, prob, radius]
         def step (x 10, X, W=None, h=0.2):
             n = X.shape[0]
             d = X.shape[1]
             superweight = 0. #superweight is the kernel X weight for each item
             x 11 = np.zeros((1,d))
             if W is None:
                 W = np.ones((n,1))
             else:
                 M = M
             for j in range(n):
                 kernel = kernelize(x 10, X[j], h, d)
                 kernel = kernel * W[j]/(h**d)
                 superweight = superweight + kernel
                 x 11 = x 11 + (kernel * X[j])
             x_11 = x_11/superweight
             density = superweight/np.sum(W)
             return [x_l1, density]
         def kernelize(x, y, h, degree):
             kernel = np.exp(-(np.linalg.norm(x-y)/h)**2./2.)/((2.*np.pi)**(degree/2))
             return kernel
         def density(X,D,h,degree):
             sum1=0
             for i in range(D.shape[0]):
                 k=kernelize(X,D[i],h,degree)
                 sum1=sum1+k
             d = 1./D.shape[0]/h**degree*sum1
             return d
         def DENCLUE(D.h.xi.eps):
```

```
In [32]: class DENCLUE (BaseEstimator, ClusterMixin):
             def init (self, h=0.2, eps=0.0001, min density=0., metric='euclidean'):
                 self.h = h
                 self.eps = eps
                 self.min density = min density
                 self.metric = metric
             def fit(self, X, y=None, sample weight=None):
                 if not self.eps > 0.0:
                     raise ValueError ("eps must be positive.")
                 self.n samples = X.shape[0]
                 self.n features = X.shape[1]
                 density attractors = np.zeros((self.n samples, self.n features))
                 radii = np.zeros((self.n samples,1))
                 density = np.zeros((self.n samples,1))
                 #create default values
                 if self.h is None:
                    self.h = np.std(X)/5
                 if sample weight is None:
                     sample weight = np.ones((self.n samples,1))
                     sample weight = sample weight
                 #initialize all labels to noise
                 labels = -np.ones(X.shape[0])
                 #climb each hill
                 for i in range(self.n samples):
                     density attractors[i], density[i], radii[i] = hill climb(X[i], X, W=sa
         mple weight,
                                                               h=self.h, eps=self.eps)
                 #initialize cluster graph to finalize clusters. Networkx graph is
                 #used to verify clusters, which are connected components of the
                 #graph. Edges are defined as density attractors being in the same
                 #neighborhood as defined by our radii for each attractor.
                 cluster info = {}
                 num clusters = 0
                 cluster info[num clusters]={'instances': [0],
                                              'centroid': np.atleast 2d(density attractors
         [0])}
                 g clusters = nx.Graph()
                 for j1 in range(self.n samples):
                     g clusters.add node(j1, attr dict={'attractor':density attractors[j1],
         'radius':radii[j1],
                                          'density':density[j1]})
                 #populate cluster graph
                 for j1 in range(self.n samples):
                     for j2 in (x for x in range(self.n samples) if x != j1):
                         if g_clusters.has_edge(j1,j2):
                             continue
                         diff = np.linalg.norm(g clusters.node[j1]['attractor']-g clusters.n
         ode[j2]['attractor'])
                         if diff <= (g clusters.node[j1]['radius']+g clusters.node[j1]['radi</pre>
         us']):
                             g_clusters.add_edge(j1, j2)
                 #connected components represent a cluster
                 clusters = list(nx.connected_component_subgraphs(g_clusters))
                 num_clusters = 0
```

```
In [39]: ## Main function

# reading the file
iris = pd.read_csv(FILE_NAME, header=None)

data = iris
data = np.array(data)
samples = np.mat(data[:,0:4])
true_labels=data[:,-1]
labels=list(set(true_labels))
true_ID=np.zeros((3,50))
index=range(len(true_labels));
for i in range(len(labels)):
    true_ID[i]=[j for j in index if true_labels[j]==labels[i]]

d = DENCLUE(0.2, 0.0001)

print(d)

DENCLUE(eps=0.0001, h=0.2, metric='euclidean', min density=0.0)
```

22...22. (epo 0.0001, 11 0.12, 1100110 0001110011 , 11111_0001101 , 11111_0001101)

```
In []:
```