## Carbon

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## 2.Clustering

## 1. Load data

```
#load module data
import data
#create a dataset object and read data from file sample.txt
sample = data.DataSet()
#To read nominal data, you have to add argument 'nominal',
default is 'numeric'
#Read data from 'sample.txt'
sample.read('sample.txt', 'numeric')
#create train dataset and test dataset using 1:10 hold out
train,test = data.holdOut(sample,0.1)
```

## 2. kMeans

A kMeans algorithm which also support bi-kMeans.

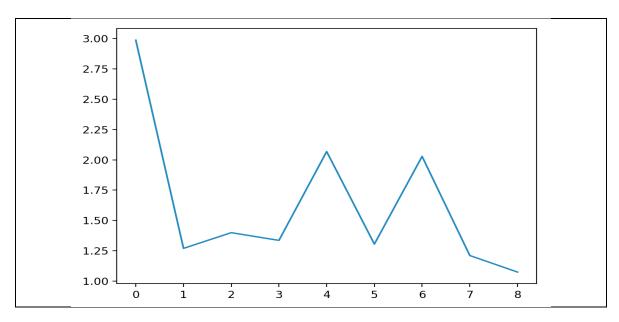
The algorithm only works for: numerical data

Parameters:

k: int from 1 to inf dist: 'euclidean'

method: 'KMeans', 'biKMeans'

```
#load kMeans module
from clustering import kMeans
#create an instance with k=3, euclidean distance, KMeans
instance = kMeans.build(k=3, dist='euclidean', method='KMeans')
#train the classifier with train data and k=4
centroids,clusters = instance.cluster(train.x)
#centroids contains the centers of each cluster
centroids
matrix([[ 7.25555556, 4.87777778, 5.12222222, 4.9
4.02222222, 9.04444444, 5.27777778, 3.75555556, 1.7
[ 2.96933962, 1.26650943, 1.4009434 , 1.31603774,
2.0754717 , 1.29009434, 2.02830189, 1.22169811, 1.07075472],
        [7.04310345, 8.40517241, 8.05172414, 6.72413793, 6.5]
, 7.17241379, 6.89655172, 7.88793103, 3.3362069 ]])
#clusters contains the cluster labels and distances from the
centers
clusters
matrix([[
            1.
                    , 5.854769271,
           1. , 6.9962787 J,
2. , 125.31599287 J,
                     , 49.505648041,
            2.
            2.
                         62.022889421,
                         53.57461356]])
           2.
#to view the clusters by plot 2 features
kMeans.view(train, centroids, clusters)
```



```
#fast start
import data
sample = data.DataSet()
sample.read('sample.txt')
train,test = data.holdOut(sample,0.1)
from imp import reload

reload(CART)
clf = CART.build()
clf.train(train, tolS=1,tolN=4,model=False)
clf.classify([5, 4, 4, 5, 7, 10, 3, 2, 1])
clf.view('Uniformity of Cell Size', 'Uniformity of Cell Shape',train)
```