

A K Means Based Genetic Algorithm for Data Clustering

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Overview

- 1 K Means Clustering
- 2 Evolutionary K Means Clustering
- 3 Proposed Method (CluGA)
- 4 Conclusion

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K Means Clustering

- Unsupervised data analysis technique in categorizing data
- K-means: most widely used clustering algorithms
- K-means involves following major steps:
 - 'n' number of centroid is selected (equal to the number of clusters)
 - Objects are assigned to the cluster
 - Centroid is updated and 2 and 3 are repeated until no objects changes its membership to a cluster
- Number of clusters should be preassigned

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Evolutionary K Means Clustering

- Genetic K means algorithm, Krishna K, Murty MN, 1996
- Capable of exploring search space and escaping from global minima
- Prior knowledge of number of clusters is not required
- High values of the evaluation indexes

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CluGA(Proposed Method)

- Introduction
- Encoding
- Initialization
- Crossover
- Mutation
- Fitness Function
- Results

Introduction

- Consists GA integrated with K Means principle
- Each individual is a string of length equal to number of dataset objects $[2, \dots, k_{max}]$
- Crossover operator obtains the offspring by adopting a group based strategy
- Mutation operator reassigns an object x belonging to a cluster C_i to another cluster C_j if x is closer to C_j .

- Uses Label based encoding
- Each object is in a position i of the chromosome and also associated with an integer in the alphabet $[2, \dots, k_{max}]$
- Renumbering procedure is used after application of genetic operator.
- Improves efficiency and avoids the presence of different string representing same solution

Initialization

- Uses approach used by $KMeans_{++}$
- $KMeans_{++}$ has different way of selecting center compared to regular K-Means
- $KMeans_{++}$ has following steps:
 - Choose the first cluster center $c_1 = x$ uniformly at random from the dataset X
 - Choose other cluster from remaining data objects
 - Repeat 2 until the k_{max} cluster has been found
- Has less convergence time, high speed and better accuracy

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Project Goal

- Build a binary classifier to detect melanoma in images of skin lesions using deep CNN models
- Use images within the same patient
- Use patient-level contextual information (metadata)
- AUC evaluation metric

THANK YOU