

A new grouping genetic algorithm for clustering problems

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WINTER 2017



Introduction

- Clustering is an important subgroup of unsupervised learning techniques consisting in grouping data objects into disjoint groups of clusters.
- Uses include pattern recognition, bio-engineering, image quantization, renewable energy prediction, etc.
- Evolutionary computing algorithms (EAs) have been widely applied to clustering problems due to their capacity to be applied to very different problems with very few changes.

Clustering evaluation

- Validation or evaluation of the resulting clustering allows analyzing the result in terms of objective measures.
- Two groups of evaluation methods
 - Supervised measures :
 - Rand index (R)
 - Jaccardindex(J)
 - Unsupervised measures :
 - Davis-Bouldin Index (DB)
 - Silhouette coefficient(S)

Proposed grouping genetic algorithm

- GGA is a class of evolutionary algorithm especially modified to tackle grouping problems, i.e. problems in which a number of items must be assigned to a set of predefined groups. (by Falkenauer)
- Problem encoding:
 - Separating each individual in the algorithm into two parts : $c = [l|g]$

$$l1, l2, \dots, lN | g1, g2, \dots, gk$$

example :

1 3 2 1 4 1 1 2 3 2 1 3 4 2 1 | 1 2 3 4



Fitness Function

- Two different fitness evaluations
 - Davis-Bouldin Index (DB):

$$DB(U) = \frac{1}{k} \sum_{i=1}^k \max_{i \neq j} \left\{ \frac{\sum_{x \in C_i} d^2(\mathbf{x}, \mu_i) + \sum_{x \in C_j} d^2(\mathbf{x}, \mu_j)}{d^2(\mu_i, \mu_j)} \right\}$$

- Silhouette coefficient (S):

$$s_j = \frac{a_j - b_j}{\max(a_j, b_j)},$$

$$S_j = \sum_{\mathbf{x}_j \in C_j} s_j,$$

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Selection operator

- Rank-based wheel selection mechanism

- $$f = \frac{2 \cdot R}{\varepsilon \cdot (\varepsilon + 1)}$$

- Static : Probabilities of survival (given by f) do not depend on the generation, but on the position of the individual in the list.

Crossover operator

- The probability of crossover must be high in the first stages of the algorithm, and moderate in the last ones in order to properly explore the search space.

$$P(j) = P_i - \frac{j}{TG} (P_i - P_f)$$

- (a) ind 1=[1 3 2 1 4 1 1 2 3 2 1 3 4 2 1 | 1 2 3 4]
ind 2=[3 1 2 1 3 2 2 1 3 1 2 3 2 2 2 | 1 2 3]
- (b) offspring=[- 3 2 - - - 2 3 2 - 3 - 2 - | 2 3]
- (c) offspring=[- 3 2 1' - 2' 2' 2 3 2 2' 3 - 2 2' | 2 3 1' 2']
- (d) offspring=[3 3 2 1' 1' 2' 2' 2 3 2 2' 3 2 2 2' | 2 3 1' 2']
- (e) offspring=[2 2 1 3 3 4 4 1 2 1 4 2 1 1 4 | 1 2 3 4]

Mutation operator

- Mutation operator includes small modifications in each individual of the population with a low probability, in order to explore new regions of the search space and also scape from local optima.
- Two different mutation operators :
 - Mutation by cluster splitting :

[2 2 1 3 3 4 4 1 2 1 4 2 1 1 4 | 1 2 3 4] → 2 2 1 3 3 4 4 5 2 1 4 2 5 1 4 | 1 2 3 4 5

- Mutation by clusters merging :

[2 2 1 3 3 4 4 1 2 1 4 2 1 1 4 | 1 2 3 4] → 2 2 1 3 3 2 2 1 2 1 2 2 1 1 2 | 1 2 3

$$P(j) = P_i + \frac{j}{TG} (P_f - P_i)$$

Replacement and elitism

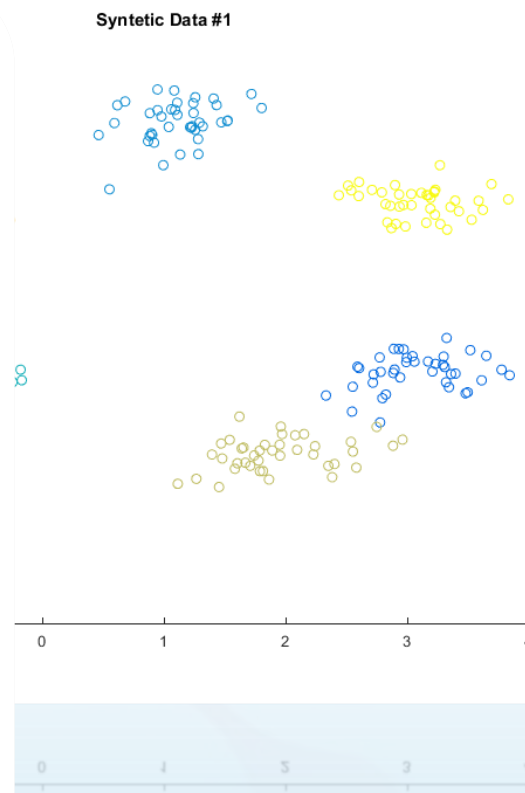
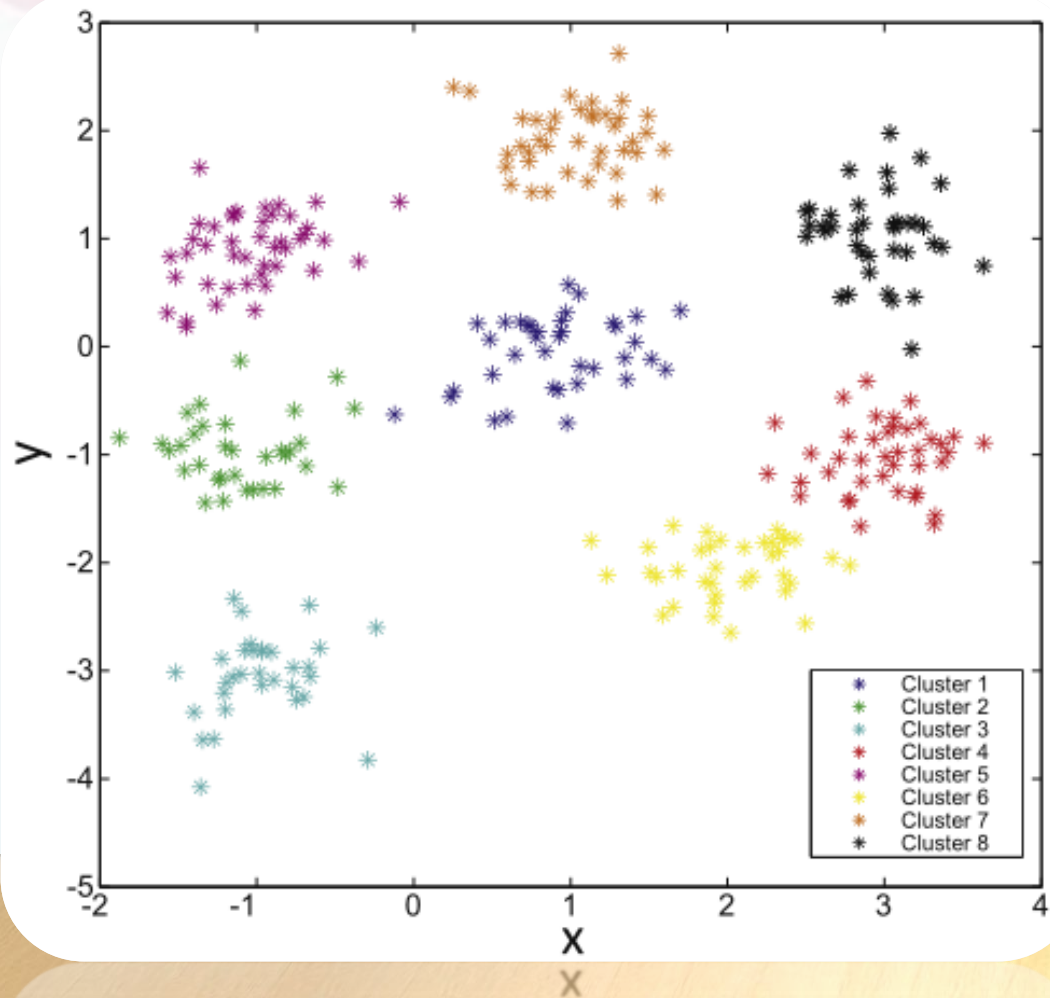
- Elitist schema is also applied , the best individual in generation j is automatically passed onto the population of generation $j + 1$.
- Best solution encountered so far in the evolution is always kept by the algorithm.
- Other individuals of population is formed by children.



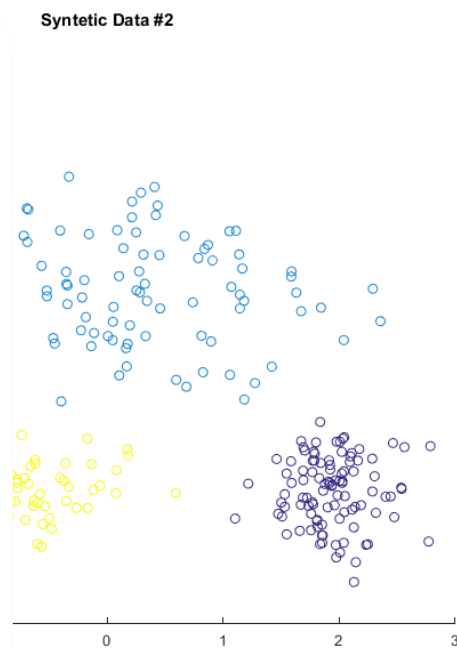
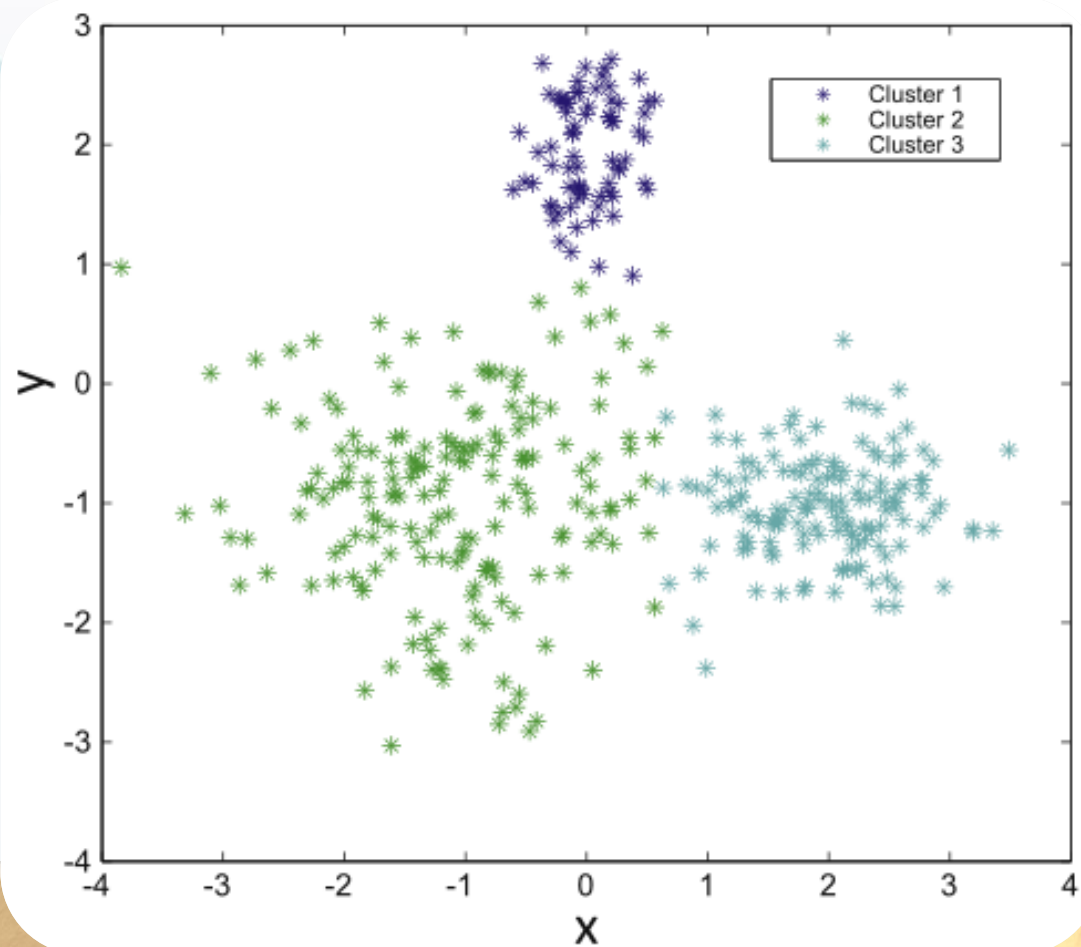
Local search

- Local search procedure tries to find local optimums in a close neighborhood of an individual.
- The implemented local search works over the element section of the individuals.
- For each observation, this operator determines the objective function variation obtained when the observation is assigned to the other clusters in the solution.
- keep the assignment with the largest objective function.
- Time consuming , so small probability.

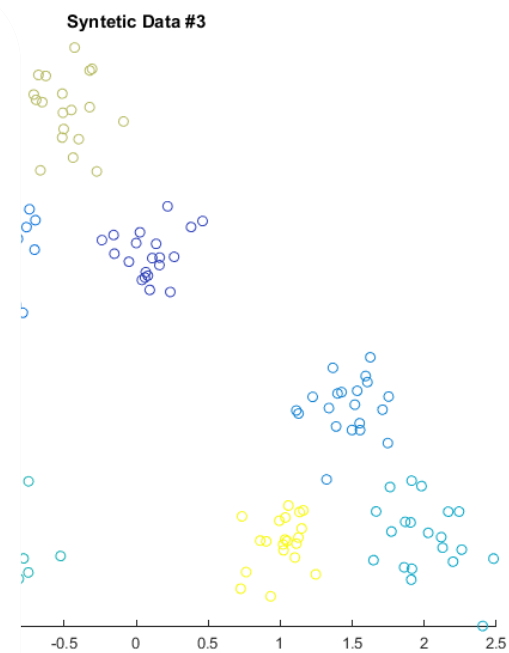
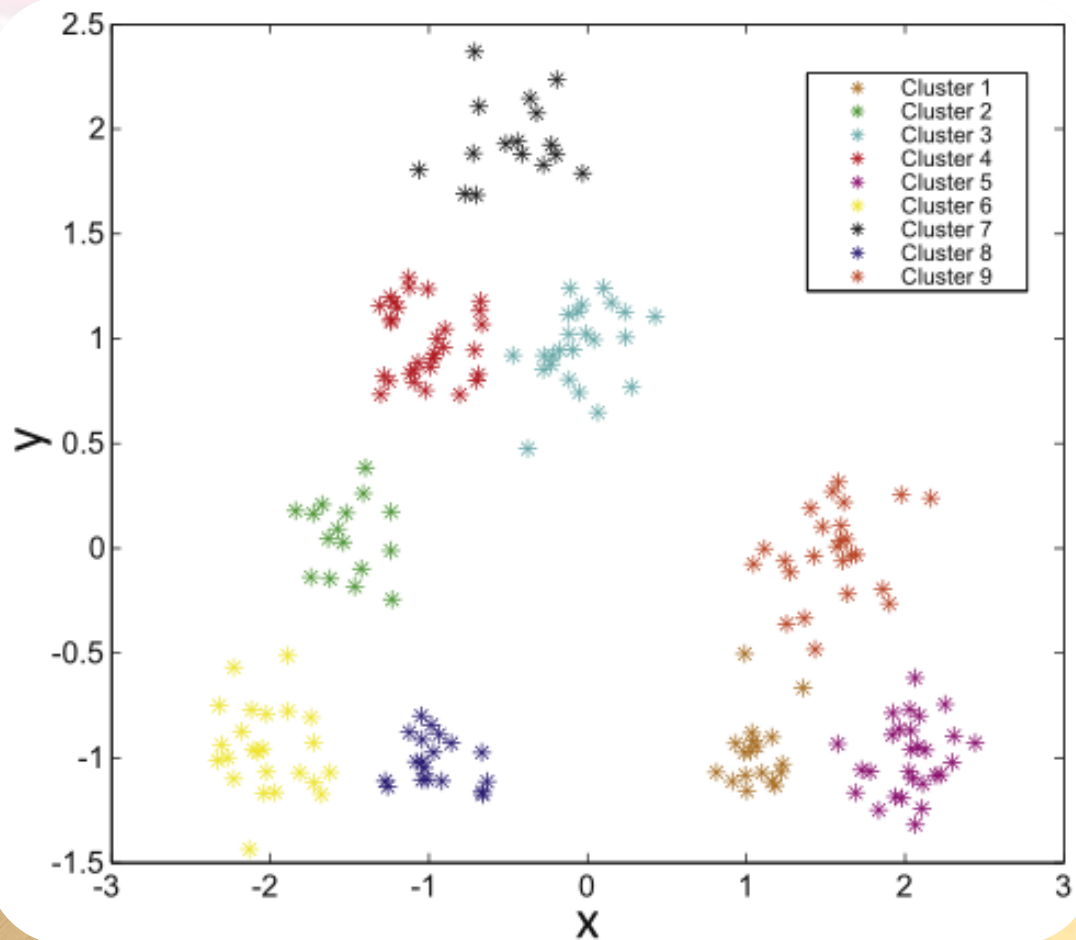
Experiments and results



Experiments and results



Experiments and results



Experiments and results : Iris Dataset

Our Results

Fitness function	# Clusters	Best Fitness	RI
DB index	3	1.192004	0.87964
S index	3	-0.654043	0.874809

Paper Results

Fitness function	# Clusters	RI
DB index	3	0.8731
S index	3	0.8995

Experiments and results : Wine Dataset

Our Results

Fitness function	# Clusters	Best Fitness	RI
DB index	3	1.6248441	0.72424
S index	3	-0.656109	0.744239

Paper Results

Fitness function	# Clusters	RI
DB index	3	0.7310
S index	3	0.7220

References :

- Brown, E. C., & Sumichrast, R. T. (2005). Evaluating performance advantages of grouping genetic algorithms. *Engineering Applications of Artificial Intelligence*, 18(1), 1-12.
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