

A Fixed-Length Subset Genetic Algorithm for the p-Median Problem

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Abstract. In this paper, we review some classical recombination operations and devise new heuristic recombinations for the fixed-length subset. Our experimental results on the classical p-median problem indicate that our method is superior and very close to the optimal solution.

1 Fixed-Length Subset Recombinations

We study the Fixed Length Subset Genetic Algorithm (*FLS-GA*), whose candidate solutions are represented by the fixed-length subset (*FLS*), which can be defined as any subset with a fixed size for a given set. In FLS-GA, we adopt a *subset encoding* [CHWS97], which uses a list of elements to represent the candidate FLS. [Rad93] studies two *pure recombinations* for FLS, which are Random Respectful Recombination (*RRR*) and Random Assorting Recombination (*RAR*). We extend them to *heuristic recombinations*.

1. Construct candidate set S' , and inherited pattern s_0 , from FLSs A and B ;
2. Choose sub-optimal FLS from S' using the heuristic procedure $H(S', s_0)$.
Let the result of $H(S', s_0)$ be the child of recombinations;

For FLS-GA, we could use RRR and RAR respectively to construct candidate set S' , and inherited pattern s_0 , leading the following two heuristic recombinations.

– Heuristic RRR (*H-RRR*)

The inherited pattern $s_0 = A \cap B$ and the candidate set $S' = A \cup B - A \cap B$. Thus the size of the inherited pattern $|s_0|$ is equal to $|A \cap B|$, while the size of the candidate set $|S'|$ is $|A \cup B| - |A \cap B|$;

– Heuristic RAR (*H-RAR*)

Each element of the inherited pattern s_0 , is chosen from s_0 with probability q . And each element of the candidate set S' , is chosen from $A \cap B - s_0$ with probability p_0 , or from $A \cup B - A \cap B$ with probability p_1 , or from $S - A \cup B$ with probability p_2 .

Moreover, to balance the diversity and pattern, an *adaptive heuristic recombinations* can be designed as follows.

– Threshold H-RAR (*T-H-RAR*)

Different heuristic recombinations are used in different situations of diversity.

1. If the diversity of the current population is less than a threshold H , we adopt H-RAR with a smaller q and bigger p_2 to increase the diversity;
2. If the diversity of the current population is larger than a threshold H , we adopt H-RAR with a bigger q and smaller p_2 to decrease the diversity;

2 FLS-GA Application to P-Median Problem (PMP)

PMP [ODE03] can be formulated as a FLS optimization problem.

– Instance:

1. A set $S = \{v_1, \dots, v_m\}$ with m vertices;
2. A positive number $p \leq m$;
3. A distance matrix $D_{m \times m}$, where D_{ij} represents the distance between vertex v_i and vertex v_j ;
4. A function $F(s) = \sum_{\forall v_i \in S} (\min_{\forall v_j \in s} D_{ij})$, $\forall s \subseteq S$;

– Constraint: $|s| = m$

– Output: A fixed length subset s^* , where $s^* \subseteq S$ and $|s^*| = m$.

– Objective: To minimize the value of $F(s^*)$;

Two heuristic procedures are adopted for when we applying FLS-GA to PMP.

– Decreasing Heuristics (*DH*)

For a given candidate set S' and an inherited pattern s_0 , firstly add all elements of S to the result FLS. Then select the rest $(p - |s_0|)$ elements from $S' - s_0$ one by one. In each round, the element, which decreases the objective function most, is added to the result FLS.

– Increasing Heuristics (*IH*) [ODE03]

For a given candidate set S' and an inherited pattern s_0 , delete $(|S'| - p)$ elements from S' one by one. In each round, the element, which is excluded from s_0 and increases the objective function the least, is deleted from S' .

The experimental results have shown that the FLS-GAs with heuristic recombinations over-perform the GA with pure genetic operations, and that the hybrid heuristic genetic recombinations exhibit the best results over all. For most problems, FLS-GAs with T-H-RAR can obtain solutions that are very close to the optimal solution (the gap $< 0.1\%$).

References

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