An Immune Operation for Lethal Chromosomes of Genetic Algorithm

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Abstract: This paper proposes an immune operation to revive the lethal chromosome (LC) for genetic algorithm (GA) so that LCs can be utilized instead of being abandoned. In general GA, LCs correspond to the infeasible solution and are eliminated from the population. However, the LCs with evolutions contain some excellent building block, we can reuse them instead of wasting resources. By this way the performance of GA is improved availably.

Keywords: genetic algorithm, lethal chromosome, artificial immune, MKP

1 Introduction

In a population pool of GA, due to crossover and mutation operations, sometimes, LCs are produced at high rate, especially to the combinatorial optimization problems having rigorous constraint. The more number of LCs increase in the population, the worse searching performance of GA decreases. At the worst case, algorithm would stop. Researches focusing on problem of LCs are still rare. Iima Hitoshi^[1] (1995) had investigated the effects of LC to performance of GA, but hadn't supplied a method to deal with. The general method to LCs is eliminating them from its population. In this paper we propose a method to revive and utilize the LCs for GA based on artificial immune theory, which combining the evolutional information of the chromosome with the characteristic information of the problem. We apply the proposed algorithm to one of the typical constrained optimization problem, Multidimensional Knapsack Problems (MKP).

2 Multidimensional Knapsack Problems

Let us explain the Multidimensional knapsack problem (MKP) briefly. MKP is a well-known NPhard problem which is formulated as:

$$Maximize \sum_{j=1}^{n} v_{j} x_{j}$$
 (1)

s.t.
$$\sum_{j=1}^{n} w_{ij} x_{j} \le c_{i} \quad (i = 1 \sim m)$$
 (2)
 $x_{i} \in \{0,1\} \quad (j = 1 \sim n),$

where n is the number of objects and m is the number of resources, v_i is the value associated with object j, w_{ij} is the consumption of resource i for object j, c_i is the available quantity of resource i (capacity of knapsacks for i^{th} resource), and x_i is the decision variable with object j and is set to 1 if j is selected (and set to 0

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otherwise). Constraints c_i (i=1,...,m) described in Eq.(2) are called knapsack constraints, so the MKP is also called the m-dimensional knapsack problem.

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In order to utilize the LCs instead of abandoning them, we extract the excellent building block from the LC firstly, and then restructure a now chromosome based on extracted building block with immune operation, which is composed of vaccine and vaccination according to immune theory.

3.1 Active ingredients of LC

The LCs with evolutions contain some active ingredients which are similar to the parts of the optimal solution; we call them excellent building blocks. To extract excellent building block from LCs, we give a method to estimate the building block, and with it extract a better one according to estimate from n/2 to n to length of building block.

For building block of LC $l_1l_2...l_n$, a binary string $f_1f_2...f_n$ is used to mark genes which belongs to building block, where $\forall i \in (1,2, ..., n), f_i=1$ denotes l_i belongs to building block, $f_i=0$ denotes not. The estimate of building block is defined as following:

$$e(f_1 f_2 ... f_n) = \frac{v(I - u)}{h}$$
 (3)

$$e(f_{1}f_{2}...f_{n}) = \frac{v(l-u)}{k}$$
where $v = \sum_{j=1}^{n} f_{j}v_{j}l_{j}, u = \max_{i} (\frac{\frac{n}{k}\sum_{j=1}^{n} f_{j}w_{ij}l_{j} - c_{i}}{c_{i}}), v \text{ is }$

different from fitness of chromosome, which is parts of fitness, and is not bounded by c_i of knapsack. k is number of genes in building block marked by $f_1f_2...f_n$ in LC, it is also called length of building block.

The steps of extracting the excellent building block from LC are outlined as following:

1: *d*←0;

2: for j=n/2 to n do:

3: $e_1e_2...e_n \leftarrow (0,0,...,0)$;

select j genes randomly from $l_1 l_2 ... l_n$ and set corresponding $e_k \leftarrow 1$;

5: **if** $(e(e_1e_2...e_n)>d)$ then

6: $f_1f_2...f_n \leftarrow e_1e_2...e_n;$

7: $d \leftarrow e(e_1e_2...e_n);$

8: end if

9: end for

10: get $f_1f_2...f_n$ to mark the excellent extracted building block in $l_1l_2...l_n$;

3.2 Training vaccine

In proposed IGA, with extracting excellent building block, we introduce the immune theory into GA to

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revive the LCs created by genetic operation.

In the beginning of evolution of GA, construct a schema of vaccine: $s_1s_2...s_n$, and set $s_i=0$ (i=1,...,n). During of evolution, when each LC $l_1l_2...l_n$ appears due to operation of GA, make following operation:

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if l_i = 1, add 1 to s_i; (i=1,...,n) otherwise, s_i is unchanged; (i=1,...,n)
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so $s_i(i=1,...,n)$ will increase with evolution process forward, and s_i , $s_j(i \neq j)$ will usually being different each other. For convenience to state, name the schema of vaccine $s_1s_2...s_n$ as vaccine.1.

In the other hand, we construct another schema of vaccine $b_1b_2...b_n$, to record excellent genes tendency. In the course of GA evolution, there is always one of the best chromosomes is retained; the old best chromosome is replaced when a new one appears. With setting $b_1b_2...b_n \leftarrow (0,0,...,0)$, during of evolution, whenever a new best chromosome $l_1l_2...l_n$ appears, make following operation:

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if l_i = 1, add \hat{1} to b_i; (i=1,...,n) otherwise, b_i is unchanged; (i=1,...,n)
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so $b_i(i=1,...,n)$ will increase with evolution process forward too, and b_i , $b_j(i \neq j)$ will usually being different each other. We call the $b_1b_2...b_n$ as vaccine.2.

3.3 Vaccination

To a worthwhile LC we have much available resources to deal with it, excellent building block, *vaccine.1* and *vaccine.2*, to revive it, there are two operations, and we summarize them in the following.

(1) The first operation is called immune.1, which sets the remaining genes excepted the genes of extracted building block from LC as 1 firstly, and then according to tendency of vaccine.1, with roulette method, to select and set genes as 0 one by one until LC being revived. Explaining with pseudo-code as follows:

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1: set the LC for vaccination as l_1l_2...l_n;
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2: extract the excellent building block from LC and mark them as $f_1f_2...f_n$;

3: for i=1 to n do:

4: $if(f_i=0) l_i \leftarrow 1$;

5: end for

6: while $(l_1 l_2 ... l_n \text{ is still lethal})$ do:

7: select one l_k by roulette according to the value of s_k in the vaccine. $l_k s_1 s_2 \dots s_n$ from all $l_i = l$ with $l_i = 0$;

8: Set the selected $l_k \leftarrow 0$;

9: end while;

(2) Another operation, immune.2, sets remaining genes of LC excepted the genes of extracted building block as 0 firstly, when the chromosome become to non-lethal, according to tendency of vaccine.2, with roulette method, to select and set genes as 1 one by one until LC become to lethal again from non-lethal state, and then make revocation of the last setting of gene.

Here we explain the algorithm just by the example of MKP; to other optimal problem, ideas and methods

are similarly.

4 Experiments on MKP

We performed the computer experiments of proposed genetic algorithm with immune operations (IGA) with taking instances of MKP from the OR-Library, and compared with the experimental results of SGA (GA without utilizing the LCs). Ordinarily the evolutional curve of GA is plotted against to the number of generation of GA. In this case since IGA has larger time-complexity for a generation than SGA, we have set terminating CPU-time and plotted the evolutional curves against to CPU-time instead of generation to compare the results of IGA and SGA and to examine their performance.

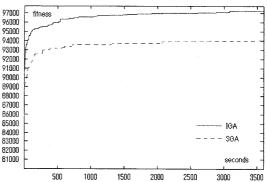


Fig.1 evolutional curve against to CPU-time

With setting population of GA as 50, termination CPU-time=3600(seconds), running times as 10, Fig.1 shows an average of evolutional curve for MKP that m=30, n=500, in which SGA finished 630.5 generations averagely and IGA is 566.4. Although SGA is slight faster than IGA in generations, IGA is rapider in CPU-time and obtains a better solution finally.

5 Conclusions

This paper proposes a method for reviving LCs replacing the present means based on immune operations. Appling it to MKP indicates that, to some cases, especially to large scale problems, IGA is more effective to deal with the LCs than SGA, this improves obviously the search performance of GA in solving the constraint optimization problems.

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