# Cross-Generational Elitist Selection SSEの収束特性について

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確率的スキーマどん欲法には、速い収束速度と制御パラメータが少ないという特徴がある。その方で、しばしば局所解に収束してしまう欠点がある。この欠点を解決するために、本研究では cross-generational elitist selection SSE (cSSE) を提案する。この方法では、cross-generational elitist selection を利用することで、SSE の個体集団の多様性を改善し、大域的探索性能を改善する。数値例において、cSSE を minimum generation gap (MGG) を用いた遺伝的アルゴリズムや Bayesian Optimization Algorithm (BOA)、SSE と比較する。計算結果により、cSSE は他のアルゴリズムよりも、短い計算時間で同程度以上の精度の解を求められることを示す。

## Convergence Analysis of Cross-Generational Elitist Selection SSE

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Stochastic Schemata Exploiter (SSE) has interesting features such as very fast convergence speed and only one control parameter. It sometimes converges to local optimum solution. In this paper, one describes cross-generational elitist selection SSE (cSSE). Since the use of the cross-generational elitist selection enhances the diversity of the individuals in the population, the global search performance is improved. In the numerical examples, cSSE is compared with genetic algorithm with minimum generation gap (MGG), Bayesian Optimization Algorithm (BOA), and original SSE. The results show that cSSE can find better solutions at shorter CPU time than the other algorithms.

## 1. Introduction

The search performance of the simple Genetic Algorithm (SGA) depends on the early convergence and evolutionary stagnation<sup>1),2)</sup>. The early convergence means that all individuals gather to same local optimum solutions at early generation and therefore, the global optimum solution can not be found. The evolutionary stagnation means that the convergence speed slows down. For overcoming these problems, a new generational alternation model, Minimal Generation Gap (MGG), was presented by Sato et.al.<sup>3)</sup>.

Bayesian Optimization Algorithm (BOA) is also one of evolutionary computations<sup>4</sup>. BOA searches a solution by using Bayesian network learned from the information of the better solutions in the population. Since offspring are generated from the stochastic model, evolutionary operations in SGA

Stochastic Schemata Exploiter (SSE) was presented by Aizawa<sup>5)</sup>. Individuals are ranked according to the descending order of their fitness. Sub-populations are generated according to the semi-order relation of sub-populations. Common schemata are extracted from the individuals in each sub-population. New individuals are generated from the schemata. Since SSE can spread better schemata over the whole population faster than the GA, the convergence speed of SSE is also faster than that of the GA. However, SSE sometimes converges to local optimum solution.

The aim of this study is to improve the search performance of SSE without sacrificing the convergence speed. For this purpose, one presents cross-generational elitist selection SSE (cSSE), in which the cross generational elitist selection<sup>6)</sup> is introduced to the original SSE. The use of the cross-generational elitist selection enhances the diversity of the population and therefore, the global search

such as selection, crossover, and mutation are not necessary.

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performance is improved. In the numerical examples, cSSE is compared with genetic algorithm with minimum generation gap (MGG), Bayesian Optimization Algorithm(BOA), and the original SSE in some numerical examples.

#### 2. cSSE Algorithm

#### 2.1 Process of cSSE

The cSSE process is summarized as follows.

- An initial population is constructed by randomly generating M individuals.
- (2) The fitness function of individual is estimated and individual is ranked according to the descending order of their fitness.
- (3) If the criterion is satisfied, the process stops.
- (4) M sub-populations are generated according to the order of the individuals.
- (5) Common schemata are extracted from the individuals in sub-populations.
- (6) M offspring are generated from M schemata.
- (7) The cross generational elitist selection makes a new population by the individuals selected from parent population and offspring population.
- (8) A generation is incremented and the process returns to 2.

## 2.2 Defining sub-populations

Sub-populations are generated according to the semi-order relation between the subpopulations. The semi-order relation can be explained as follows.

The population P is composed of the individuals  $c_1, c_2, \cdots, c_M$ . The individuals are numbered according to the descending order of their fitness. The sub-population of the population P is referred as S. When the individual  $c_k$  is excluded from S, the new sub-population is as  $S-c_k$ . The operator  $\cup$  denotes the union of sets. The rank of the worst individual in the sub-population S is represented as L(S). Since the worst individual in the sub-population S is  $c_{L(S)}$ ,  $c_{(L(S)+1)}$  denotes the worse one by one rank than  $c_{L(S)}$  in the whole population P.

We can find the following semi-order relation among the sub-populations of the population P;

• The average fitness value of individuals in the sub-population S is higher than that in the sub-

- population  $S \cup c_{(L(S)+1)}$ .
- The average fitness value of the individuals in the sub-population S is higher than that in the sub-population (S − c<sub>L(S)</sub>) ∪ c<sub>(L(S)+1)</sub>.

For example, the semi-order relation shows the order of the schemata as follows:

- (1) Schema of  $c_1$ ; i.e, chromosome of  $c_1$  itself.
- (2) Common schema between  $c_1$  and  $c_2$ .
- (3) Schema of  $c_2$ .
- (4) Common schema between  $c_1$  and  $c_3$ .
- (5) ····

#### 2.3 Extracting Common Schemata

After defining the sub-populations, the common schemata are determined as the product set of the chromosomes of the individuals in the subpopulations.

#### 2.4 Generating New Individuals

The extracted schemata are composed of three characters; "0"s, "1"s, and "\*"s. The new individuals are defined by randomly replacing "\*" by "0" or "1".

#### 2.5 Cross Generational Elitist Selection

Algorithm of cross generational elitist selection is written as follows<sup>6</sup>).

- (1) At generation t-1, offspring are generated from individuals in the population.
- (2) Populations of parents and offspring are indicated with P(t-1) and O(t-1), respectively.
- (3) P(t-1) and O(t-1) are merged to new population P'(t-1).
- (4) Individuals in P'(t-1) are ranked according to their fitness.
- (5) The population P(t) is generated by selecting best M individuals from P'(t-1).

## 3. Numerical Examples

cSSE is compared with GA with MGG, BOA, SSE, and cSSE in some problems<sup>7)</sup>.

In GA with MGG, two-point crossover of crossover rate = 1 is employed and all individuals are replaced at every generations. In all algorithms, the best mutation rates are determined from numerical experiments.

Maximum generation is 40,000 for deception problem and 15,000 for knapsack problem, respectively. Population size is specified as  $N_i$  =

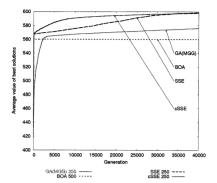


図1 だまし問題における最良解の適応度平均値

Fig. 1 Average fitness value of solutions on deception problem

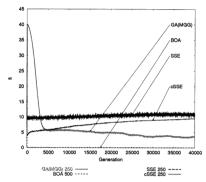


図 2 だまし問題における多様性パラメータ  $f_i$ Fig. 2 Diversity parameter  $f_i$  in deception problem

10, 50, 100 or 250 for GA with MGG, SSE and cSSE and  $n_i = 20, 100, 200$  or 500 for BOA, respectively. Since BOA replaces half population size at every generations, computational cost of fitness function is half as much as the other algorithms. For equalizing the computational cost of all algorithms, the population size of BOA is twice as many as the other algorithms. Simulations are performed 50 times from different initial populations. The average fitness values of the best individuals are compared.

#### 3.1 Deception Problem

The deception problem is defines as the summation of the 4-bit deception sub-problem<sup>7)</sup>. The objective function f is defined as

$$f_{deception} = \sum_{i=1}^{n} f_d(x_i) \tag{1}$$

where n denotes the number of 4-bit deception problem and n = 10. The design variable  $x_i$  of the problem are defined as  $x_i \in 0000, 0001, \dots, 1111$ .

The history of average fitness of the best individual is shown in Fig.1. The abscissa and the ordinate denote the generation and the fitness value, respectively. We notice from Fig.1 that the convergence speed of MGG is the slowest and that the cSSE is the fastest among them. Although BOA has very fast convergence speed, it may be attracted to a local optimum solution because the final BOA solution is worse than the other.

Next, we will discuss the diversity of solutions in deception problem. For that purpose, we will define the diversity parameter  $f_i$  as follows.

$$f_i = \frac{\sum_{i=1}^{M-1} \sum_{j=i+1}^{M} |p_i - p_j|}{{}_{M}C_2}$$
 (2)

where M and p denote population size and an individual, respectively.  $|p_i-p_j|$  denotes the hamming distance between  $p_i$  and  $p_j$ . Therefore, the diversity of the individuals in the population depends on the magnitude of the parameter  $f_i$ .

Figure 2 shows the history of the parameter  $f_i$ . While the parameter in BOA decreases to zero immediately, the parameters in the others are relatively big till the final generation. We can recognize that this is the reason why the BOA is caught to the local solution.

## 3.2 Knapsack Problem

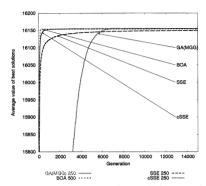
When there are n baggage in a knapsack, the knapsack problem is defined as the maximization of the value of the knapsack without exceeding the weight limit b. The problem is defined as

$$\max_{\{x_i\}} \sum_{i=1}^{n} c_i x_i$$
subject to 
$$\sum_{i=1}^{n} a_i x_i \le b$$

$$x_i \in 0, 1 \quad (i = 1, \dots, n)$$
(3)

where  $a_i$  and  $c_i$  denote the weight and the value of the baggage i, respectively. They are randomly taken within  $1 \le a_i, c_i \le 100$  and b = 10000 and n = 400.

The history of the average fitness value of the best individuals is shown in Fig.3. Figure 3 illustrates that the convergence speed of MGG is the



**図3** ナップザック問題における各世代の最適解の平均適応度 Fig. 3 Average fitness value of solutions on knapsack problem

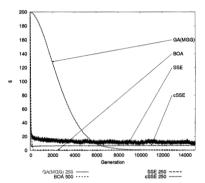


Fig. 3 Diversity parameter  $f_i$  in knapsack problem

slowest among them.

Figure 3.2 shows the history of the parameter  $f_i$ . Also in this case, the parameter in BOA decreases to zero immediately. On the other hand, the parameters in the others are relatively big till the final generation.

### 4. Conclusions

In this study, we described the cross generational elitist selection SSE (cSSE) in which the generation alternation model of original SSE was changed to cross generational elitist selection model. The presented algorithm was compared with GA with MGG, BOA, and SSE.

Numerical results shows the following features of cSSE.

In comparing search performance of algorithms, note that cSSE can find slightly or much better solution than the others in all examples. Specially, cSSE shows very good performance in the deception problem. In the problem, the average fitness value of the final cSSE solutions is better than that of GA with MGG by about 5%.

In comparing the convergence speed of algorithms, note that cSSE is much faster than GA with MGG and similar to BOA and original SEE in the deception. Although, in the knapsack problem, cSSE is slightly slower than BOA, their difference is small.

Finally, we can conclude that cSSE is better than the others from both viewpoints of the search performance and the convergence speed.

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