

Genetic Algorithm with Virus Infection for Finding Approximate Solution

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Abstract—Genetic Algorithm (GA) is modeling behavior of evolution in organic and known as one of method to solve Traveling Salesman Problem (TSP). However, GA obtains solution by overlaying generations because of being based on evolution in organic. Thus, it takes a long time to find approximate solution. While, Virus Theory of Evolution (VTE) can evolve by virus infection. VTE characteristic has sharing of information among same generation. If new algorithm is using both these characteristics of GA and VTE, convergence speed would be faster than GA. Thus, this study proposes Genetic Algorithm with Virus Infection (GAVI). GAVI algorithm is Virus Theory of Evolution (VTE) to be based on Genetic Algorithm (GA). We apply GAVI to TSP and confirm that GAVI obtains more effective result than GA.

that the proposed algorithm achieves better performance than the conventional GA.

TABLE I
THE NUMBER OF CITIES AND THE TOTAL ROUTE NUMBER

number of cities	total route number
3	1
4	3
8	$2.52 * 10^3$
16	$6.54 * 10^{11}$
32	$8.22 * 10^{33}$
64	$9.91 * 10^{86}$
128	$1.51 * 10^{213}$

I. INTRODUCTION

Traveling Salesman Problem (TSP) [1] is known as one of the combinatorial optimization problems. When Salesman tours all cities at once, TSP is the problem of finding minimum total between each city distance in route. Then n is the number of cities in TSP, total route number increase at rate proportional to many of the factorial of n . For example, when the number of n increases, the total route number explode in Table I. Therefore, exploring total route number needs amount of time in finding approximate solution. However we would like to obtain the approximate solution in TSP quickly. It is necessary to solve the TSP in other ways except exploring total routes. Finding approximate solution method of TSP is a variety of ways.

Genetic Algorithm (GA) [2], [3] is one of the popular method in variety of ways to solve the TSP and is studied by many researchers all over the world. GA is modeling behavior of evolution in organic. In genetic group, individuals with high evaluation tend to be many crossover. Thus, genetic group obtains high evaluation in average for repeating crossover. However, this method needs amount of time and convergence speed is slow. We would like to obtain approximate solution quickly. Thus, we need to propose the other method.

In this study, we propose Genetic Algorithm with Virus Infection (GAVI). One of the characteristics of the virus infection [4]-[6] conveys gene information among same generation at once. This characteristic seems to be useful for finding the approximate solution quickly. Therefore, we consider that the virus infection will help to solve the TSP. We carry out computer simulations for various parameter values and confirm

II. VIRUS THEORY OF EVOLUTION

Organic evolution is theory based on natural selection. In natural world, high fitness individuals organism survive, while low fitness individuals organism become extinct. Over the years, only higher fitness individuals survive. We call it Evolution. Thus, evolution need to overlay generations.

On the other hand, there is theory that Virus Theory of Evolution (VTE) [7]. This theory is based on the evolution by Lateral Gene Transfer (LGT) [8] in Virus infection. LGT is uptake of the gene that occur between other individuals and among other species. Without evolution inherited from parent cell to child cell, genes can evolve. Low fitness individuals possibly evolve into high fitness individuals in just one generation by LGT in Virus infection. Thus, we assume using VTE algorithm leads the approximate solution in less time and VTE theory is efficient in TSP.

III. GENETIC ALGORITHM WITH VIRUS INFECTION

Selection, Crossover and Mutation are the main functions of the GA. GAVI is a method of VTE algorithm in Virus infection to be based on GA. Flow chart of GAVI is shown in Fig. 1. New algorithm is Infection in Fig. 1. t is the number of repeating times.

Initialization

Initialization is random route selection. Number of random route selection is U .

Evaluation

Evaluation is defined by the following formula.

$$f_i = \frac{1}{d_i} \quad (1)$$

where d_i is total distance of each route and f_i is evaluation value. If d_i is low, f_i is high by this formula.

Selection

Route is selected with a probability of p_i . p_i is defined by the following formula.

$$p_i = \frac{f_i}{\sum_{i=1}^n f_i} \quad (2)$$

where n is the number of cities. Selection is the method of tends to be chosen high evaluation route. Low evaluation route is chosen a few. Selection works that high fitness individuals organism survive, while low fitness individuals organism become extinct.

Fulfill crossover condition

This section evaluates crossover condition. If parents is not fulfill crossover condition, crossover is not acted. When crossover condition is fulfill, crossover is acted.

Crossover

Crossover is to be mated the two routes. In this study, we apply sub tour exchange crossover. This way makes a search for sub tour of the both Parent A and Parent B in common. If it does not find sub tour in common, crossover algorithm is not acted. For example, between 1, 2, 5, 6 and 5, 1, 6, 2 are sub tour in Fig. 2. 1, 2, 5, 6 and 5, 1, 6, 2 are differ in line, however these are same class. Sub tour in 1, 2, 5, 6 can express 1, 2, 5, 6 and 6, 5, 2, 1, 5, 1, 6, 2 can express 5, 1, 6, 2 and 2, 6, 1, 5. Because two expressing are same about total route distance in Fig. 3. Thus, after crossover, four child exist.

Mutation

Mutation is to change the route with a certain probability in Fig. 5 and less likely to fall into local minimum.

Infection

We define elements of each route as Virus, elements select the best route in the obtained solution and part of each route is infected by this Virus at random. Number of element selects is fixed probability. For example, 3,5 is a virus and has infected the route of 6, 1, 3, 5, 2, 4 in Fig. 4. *Infection* part determines 1, 4 in the route. The route replace to 3, 5 1, 4. We call it *Infection*. *Infection* is incorporating partial optimum solution.

One route reset in random

If the obtained solution is same among number of s , one route in all routes is initialization in random. $O(t)$ is the obtained solution in number of t times, while $O(t-s)$ is the obtained solution previous number of s . Thus, $O(t) = O(t-s)$ shows that the obtained solution is same among number of s . We assume that this is efficient to escape local minimum.

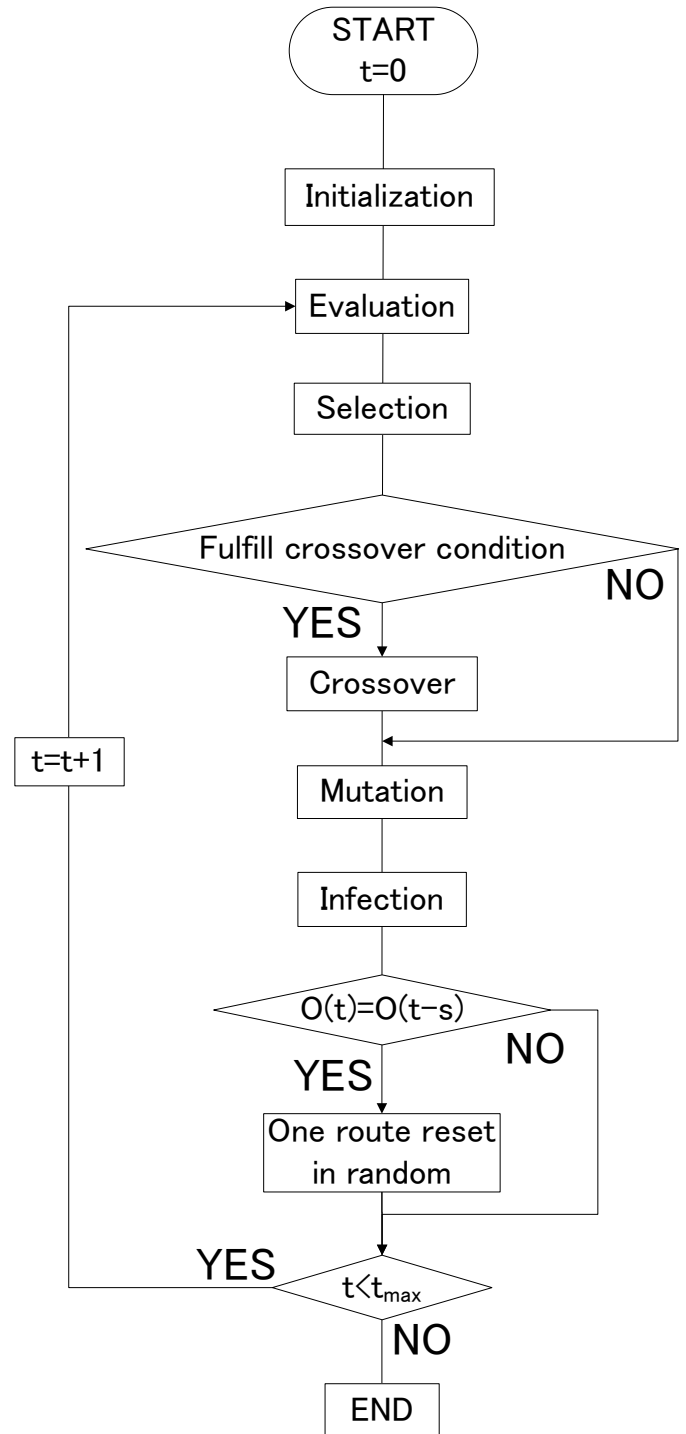


Fig. 1. Flow chart of GAVI.

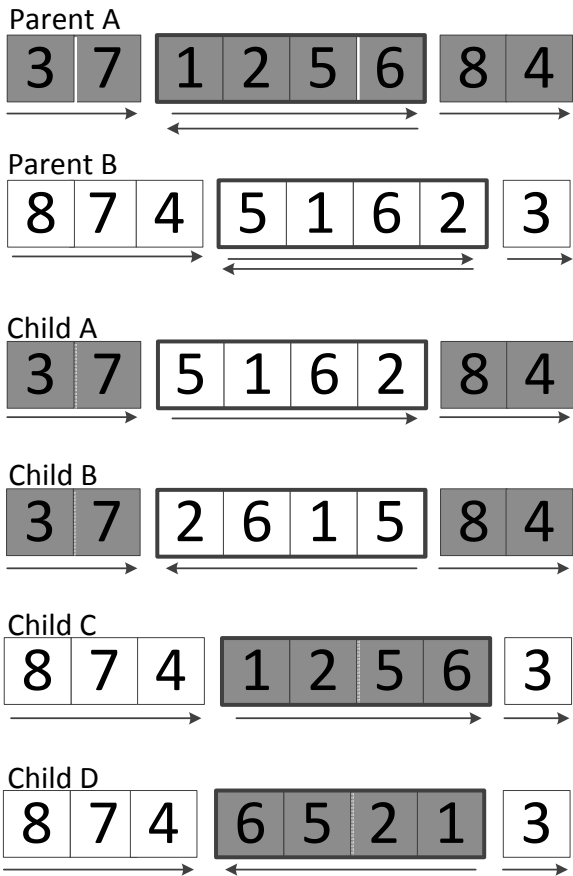


Fig. 2. The mechanism of *Infection*.

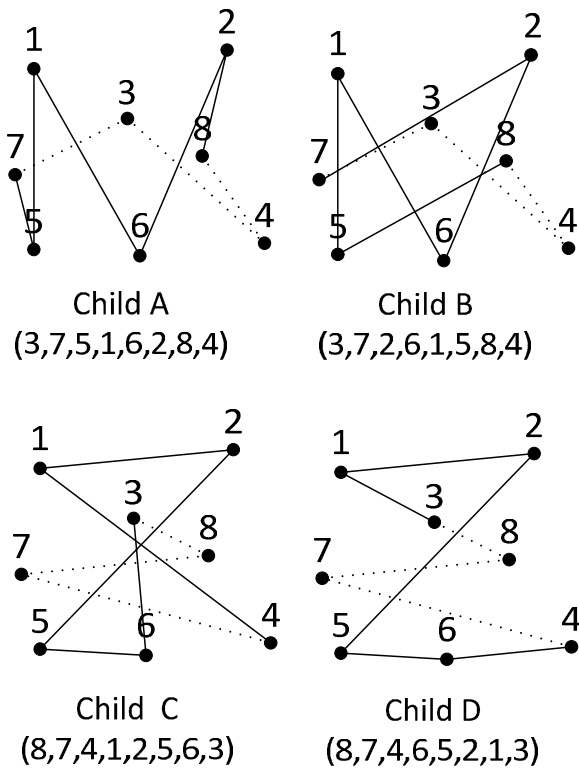


Fig. 3. The mechanism of *Infection*.

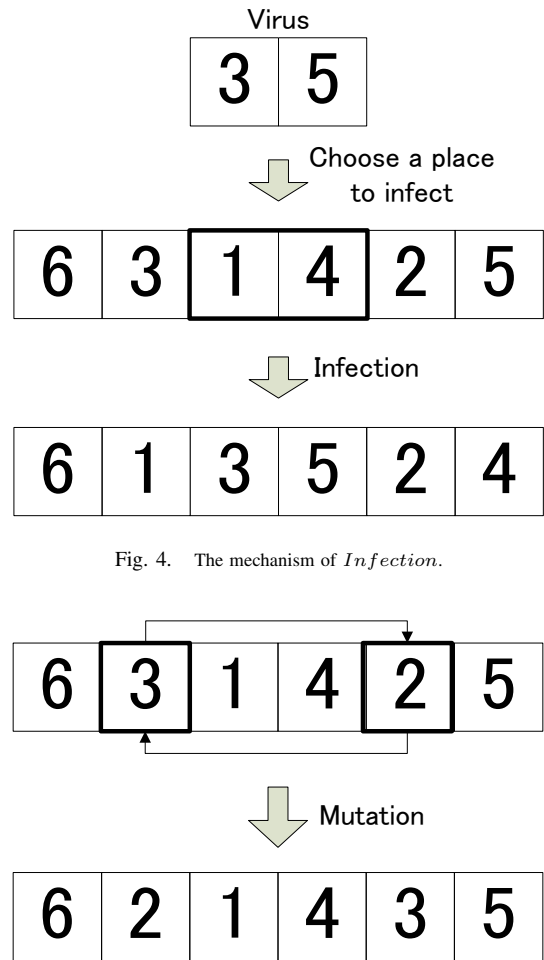


Fig. 4. The mechanism of *Infection*.

Fig. 5. The mechanism of *Mutation*.

IV. EXPERIMENTAL RESULTS

In order to compare the performance of GAVI and GA, we apply GAVI and GA to find approximate solutions in TSP such as att48 (48 cities). In this study, the number of t is 5000 times, the number of simulation is 10 times, U is 1000, number of route in and *error rate* is defined by the following formula.

$$Error\ rate[\%] = \frac{(obtain) - (optimum)}{(optimum)} \times 100 \quad (3)$$

where *obtain* shows obtained solution and *optimum* shows optimum solution. When *obtain* value approaches *optimum* value, *Error rate* is low. For example, when *obtain* value is equally *optimum* value, *Error rate* value is 0[%]. If *Error rate* value is 0[%], we would obtain optimum solution. However, *obtain* is bad solution, *error rate* value is high. In Table II, III, Ave is the average, Max is the maximum value and Min is the minimum value in the results of simulation 10 times.

Table II shows result of GA and Table III shows result of GAVI. In GAVI, s is fixed with 100 and *Mutation rate* is fixed with 0.02. GAVI can find a good solution by

Infection from result of both Table II and Table III. When *Infection rate* is 0.01, result of GAVI is the best. We assume that if *Infection rate* is high, escaping local minimum is difficult and diversity solution is loss. Thus, result of *Infection rate* = 0.1 is the better than *Infection rate* = 0.2.

Figure 6 shows the error curve of GA and Fig. 7 shows the error curve of GAVI. The number of simulation is 1 time, *Mutation rate* is fixed with 0.02 about GA and GAVI, *Infection rate* is fixed with 0.10 about GAVI. In Figure 6 and 7, Convergence speed of GAVI is faster than GA. Thus, we assume that *Infection* algorithm is efficient at first.

TABLE II
THE RESULT FOR GA IN ATT48

Algorithm type	Mutation rate	Error rate[%]		
		Max	Min	Ave
GA	0.02	6.486	0.597	2.878
	0.05	5.610	2.439	3.430
	0.10	4.621	1.280	2.096
	0.20	3.936	0.958	2.324

TABLE III
THE RESULT FOR GAVI IN ATT48

Algorithm type	Infection rate	Error ate[%]		
		Max	Min	Ave
GAVI	0.05	4.437	0.926	1.912
	0.10	2.835	0.530	1.195
	0.20	2.096	0.798	1.628

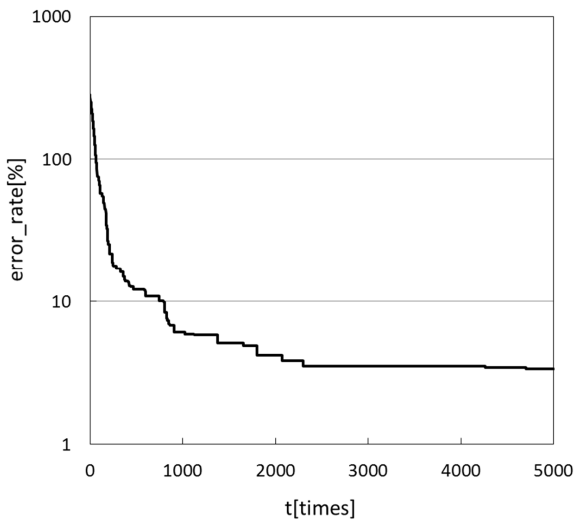


Fig. 6. error curve of GA.

V. CONCLUSION

We proposed GAVI for TSP and compared the performance of GAVI and GA to lead approximate solutions. From the

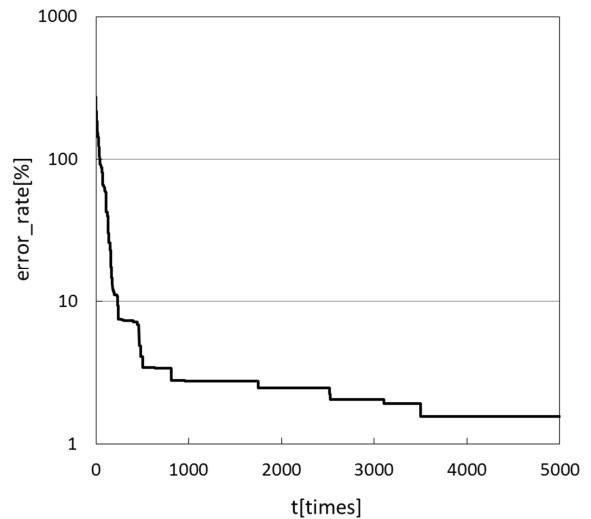


Fig. 7. error curve of GAVI.

result, the result of GAVI was better than GA. It was efficient that not only GA but also *Infection* applied in TSP. GAVI was valuable in convergence of the solution. We confirmed that VTE is efficient in TSP.

In future work, we would like to study the mechanism of *Infection* in detail. In this study, all number of element in *Infection* is depended on fixed probability. However changing fixed probability is to expected efficient. For example, large number of element is chosen frequent at first, while small number of element is chosen frequent latter. Convergence speed is fast at first, obtain solution prevent local minimum latter. Thus, we are expected to obtain a better solution in GAVI.

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