MULTI-OBJECTIVE GENETIC ALGORITHM AND ITS APPLICATIONS TO FLOWSHOP SCHEDULING

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Abstract—In this paper, we propose a multi-objective genetic algorithm and apply it to flowshop scheduling. The characteristic features of our algorithm are its selection procedure and elite preserve strategy. The selection procedure in our multi-objective genetic algorithm selects individuals for a crossover operation based on a weighted sum of multiple objective functions with variable weights. The elite preserve strategy in our algorithm uses multiple elite solutions instead of a single elite solution. That is, a certain number of individuals are selected from a tentative set of Pareto optimal solutions and inherited to the next generation as elite individuals. In order to show that our approach can handle multi-objective optimization problems with concave Pareto fronts, we apply the proposed genetic algorithm to a two-objective function optimization problem with a concave Pareto front. Last, the performance of our multi-objective genetic algorithm is examined by applying it to the flowshop scheduling problem with two objectives: to minimize the makespan and to minimize the total tardiness. We also apply our algorithm to the flowshop scheduling problem with three objectives: to minimize the makespan, to minimize the total tardiness, and to minimize the total flowtime.

I. INTRODUCTION

Flowshop scheduling problems are one of the most well known problems in the area of scheduling. The objective of minimizing the makespan is often employed as a criterion of flowshop scheduling since Johnson's work [1]. Various heuristic approaches (e.g., Dannenbring [2], Nawaz et al. [3], Osman and Potts [4] and Widmer and Hertz [5]) as well as optimization techniques (e.g., Ignall and Schrage [6] and Lomnicki [7]) have been proposed for minimizing the makespan. While these studies treated a single objective, many real-world problems involve multiple objectives. Recently several researchers have tackled multi-objective flowshop scheduling problems. For example, Ho and Chan [8] proposed a heuristic method for flowshop scheduling with bicriteria, Gangadharan et al. [9] proposed a simulated annealing heuristic for flowshop scheduling with bicriteria, and Morizawa et al. [10] proposed a complex random sampling method for multi-objective problems.

Genetic algorithms have been mainly applied to single-objective optimization problems. When we apply a single-objective genetic algorithm to a multi-objective optimization problem, multiple objective functions should be combined into a scalar fitness function. If we assign a constant weight to each of the multiple objective functions for combining them, the direction of search in the genetic algorithm is constant in the multi-dimensional objective space as shown in Fig. 1. In Fig. 1, \( f_1(\cdot) \) is an objective function to be maximized and \( f_2(\cdot) \) is to be minimized. The closed circle in Fig. 1 represents the final solution by the single-objective genetic algorithm.

Some studies have been attempted for designing multi-objective genetic algorithms since Schaffer's work [11]. Schaffer proposed the Vector Evaluated Genetic Algorithm (VEGA) for finding Pareto optimal solutions of multi-objective optimization problems. In Schaffer's VEGA, a population was divided into disjoint subpopulations, then each subpopulation was governed by its own objective function. Although Schaffer [11] reported some successful results, his approach seems to be able to find only extreme solutions on Pareto fronts as shown in Fig. 2 because its search directions are parallel to the axes of the objective space. Schaffer suggested two approaches to improve his approach in his paper [11]. One is to provide a heuristic selection preference for non-dominated individuals in each generation. The other is to crossbreed among the "species" by adding some mate selection.
Fig. 1. Direction of the search by GA with a combined fitness function.

Fig. 2. Directions of the search by Schaffer's VEGA.
problem with three objectives: to minimize the makespan, to minimize the total tardiness, and to minimize the total flowtime.

2. GENETIC OPERATIONS IN MULTI-OBJECTIVE GENETIC ALGORITHM

2.1. Selection procedure

One of the simplest methods to combine multiple objective functions into a scalar fitness solution is the following weighted sum approach (we assume that all the objective functions should be maximized):

\[
f(x) = w_1 f_1(x) + \cdots + w_i f_i(x) + \cdots + w_n f_n(x),
\]

where \( x \) is a string (i.e., solution), \( f(x) \) is a combined fitness function, \( f_i(x) \) is the \( i \)-th objective function, \( w_i \) is a constant weight for \( f_i(x) \), and \( n \) is the number of the objective functions.

If we use the weighted sum in (1) with the constant weights \( w_i \)'s, the search direction in genetic algorithms is also constant as shown in Fig. 1. Therefore we propose an idea to use variable weights in the selection procedure. The aim of this idea is to realize various search directions in Fig. 3 to search for Pareto optimal solutions. For example, we can specify variable weights as follows for a two-objective optimization problem:

\[
f(x) = w_1 f_1(x) + w_2 f_2(x),
\]

\[
\begin{align*}
    w_1 &= \frac{(i-1)}{(N_{\text{selection}} - 1)}, \quad i = 1, 2, \ldots, N_{\text{selection}}, \\
    w_2 &= 1 - w_1,
\end{align*}
\]

where \( N_{\text{selection}} \) is the number of selections in each generation. The weighting scheme in (2)-(3) means that different weights are used for selecting each pair of parent strings in our multi-objective genetic algorithm. Because \( N_{\text{selection}} \) pairs of parent strings are to be selected in each generation, \( N_{\text{selection}} \) pairs of different weights are specified in (3). In (3), the values of \( w_1 \) and \( w_2 \) are evenly distributed over the closed interval [0, 1]. In computer simulations in this paper, we used the weighting scheme in (2)-(3) for two-objective problems.

In general, we can randomly determine the value of each weight. For a multi-objective optimization problem with \( n \) objective functions (\( n \geq 2 \)), we can assign a random real number to each weight as follows when each pair of parent strings are selected for a crossover operation.

\[
w_i = \frac{\text{rnd}_i}{\sum_{j=1}^{n} \text{rnd}_j}, \quad i = 1, 2, \ldots, n,
\]

where \( \text{rnd}_i \) and \( \text{rnd}_j \) are non-negative random integers (or non-negative random real numbers). From (4), we can see that \( n \) random real numbers are generated for the weights \( w_i \)'s to calculate
the weighted sum in (1) when each pair of parent strings are selected. This procedure is iterated \( N_{\text{selection}} \) times in each generation for selecting \( N_{\text{selection}} \) pairs of parent strings for a crossover operation. In computer simulations for a three-objective problem, we used the weighting scheme in (1) and (4).

The weighted sum \( f(x) \) in (1) and (2) is used for determining the selection probability of each string in our multi-objective genetic algorithm. Because the weights \( w_i \)'s are not constant but variable, the selection probability of each string is also variable even in a single generation. This realizes various search directions in our multi-objective genetic algorithm.

2.2. Elite preserve strategy

In multi-objective optimization problems, a solution with the best value of each objective can be regarded as an elite individual. Therefore we have \( n \) elite individuals for an \( n \)-objective problem. It is natural to think that such solutions are to be preserved to the next generation in genetic algorithms.

During the execution of our multi-objective genetic algorithm, a tentative set of Pareto optimal solutions is stored and updated at every generation. We also preserve a certain number of individuals randomly selected from the tentative set of Pareto optimal solutions in addition to the \( n \) elite individuals with respect to \( n \) objectives. That is, multiple Pareto optimal solutions are used as elite individuals in our multi-objective genetic algorithm. This elite preserve strategy has an effect on keeping the variety in each population in our multi-objective genetic algorithm.

In computer simulation, three solutions were selected as elite individuals for a two-objective flowshop scheduling problem: two elite individuals with respect to two objectives and a randomly selected individual from the tentative set of Pareto optimal solution. As a variant, we can design a multi-objective genetic algorithm by randomly selecting a certain number of elite individuals from the tentative set of Pareto optimal solutions. In computer simulation for a three-objective flowshop scheduling problem, three solutions which were randomly selected from the tentative set of Pareto optimal solutions were inherited to the next generation.

2.3. Algorithm

The following genetic operations are employed to generate and handle a population (i.e., a set of strings) in our multi-objective genetic algorithm (see Fig. 4).

Fig. 4. Outline of our multi-objective genetic algorithm.
Step 0 (Initialization): Generate an initial population containing $N_{\text{pop}}$ strings where $N_{\text{pop}}$ is the number of strings in each population.

Step 1 (Evaluation): Calculate the values of the objective functions for the generated strings. Update a tentative set of Pareto optimal solutions.

Step 2 (Selection): Calculate the fitness value $f(x)$ of each string by using the weights defined as (3) or (4). Select a pair of strings from the current population according to the following selection probability. The selection probability $P(x)$ of a string $x$ in a population $\Psi$ is specified as

$$P(x) = \frac{f(x) - f_{\text{min}}(\Psi)}{\sum_{x \in \Psi} \{ f(x) - f_{\text{min}}(\Psi) \}},$$

where

$$f_{\text{min}}(\Psi) = \min\{ f(x) | x \in \Psi \}.$$  

This step is repeated $N_{\text{selection}}$ times to produce $N_{\text{pop}}$ offspring by the crossover operation in Step 3. The values of the weights $w_i$'s in (3) or (4) are specified at each of the $N_{\text{selection}}$ iterations.

Step 3 (Crossover): For each selected pair, apply a crossover operation to generate an offspring with the crossover probability $P_c$. $N_{\text{pop}}$ strings should be generated by the crossover operation ($N_{\text{selection}} = N_{\text{pop}}$ if a single offspring is generated from a pair of parent strings in the crossover operation).

Step 4 (Mutation): For each string generated by the crossover operation, apply a mutation operation with a prespecified mutation probability $P_m$.

Step 5 (Elitist strategy): Randomly remove $N_{\text{elite}}$ strings from the $N_{\text{pop}}$ strings generated by the above operations, and add the same number of strings from a tentative set of Pareto optimal solutions to the current population.

Step 6: (Termination test): If a prespecified stopping condition is not satisfied, return to Step 1.

Step 7 (User selection): The multi-objective genetic algorithm shows the final set of Pareto optimal solutions to the decision maker. A single solution (i.e., the final solution) is selected by the decision maker's preference.

In this paper, we employed the two-point crossover and the shift mutation shown in Fig. 5 for flowshop scheduling. These genetic operators were found to be good operators for a single-objective genetic algorithm for flowshop scheduling in our previous work [15].

3. SIMULATION RESULTS FOR A NUMERICAL EXAMPLE

If multi-objective optimization problems have concave Pareto fronts, weighted sum approaches with constant weights tend to fail to find entire Pareto fronts (i.e., fail to find all the Pareto optimal solutions). Our approach, however, can handle multi-objective optimization problems with concave Pareto fronts. This is shown by the following example.

Parent 1: $A\ B\ C\ D\ E\ F\ G\ H$

Child: $A\ B\ E\ D\ C\ F\ G\ H$

Parent 2: $E\ H\ A\ D\ B\ C\ G\ F$

Crossover

Mutation

Fig. 5. Genetic operators.
We applied our multi-objective genetic algorithm to a test problem with the following two objectives to be minimized:

\[ \begin{align*}
\text{Minimize } f_1(x) &= 2\sqrt{x_1}, \\
\text{Minimize } f_2(x) &= x_1(1 - x_2) + 5,
\end{align*} \tag{7}\]

subject to

\[ \begin{align*}
1 &\leq x_1 \leq 4, \\
1 &\leq x_2 \leq 2.
\end{align*} \tag{9}\]

Substituting equation (7) into equation (8), we obtained the relation between \( f_2(x) \) and \( f_1(x) \) as follows:

\[ f_2(x) = \frac{1 - x_2}{4} \left( f_1(x) \right)^2 + 5. \tag{10}\]

When \( x_2 = 2 \), equation (10) gives the Pareto front of this problem. This Pareto front forms the concave shape in the objective space as shown in Fig. 6.

In our multi-objective genetic algorithm, the fitness function \( f(x) \) was specified as

\[ f(x) = -w_1 f_1(x) - w_2 f_2(x). \tag{11}\]

Because \( f_1(x) \) and \( f_2(x) \) should be minimized, the negative sign "−" is attached to each weight in (11).

In our multi-objective genetic algorithm, the number of solutions in each population (i.e., population size) was specified as \( N_{\text{pop}} = 100 \), two point crossover was employed with the crossover probability 0.9, the mutation probability was specified as \( P_{\text{m}} = 0.01 \), and the number of elite individuals was specified as \( N_{\text{elite}} = 5 \) (five elite solutions were randomly chosen from a tentative set of Pareto optimal solutions). In Fig. 6, □ denotes the final solutions by our multi-objective genetic algorithm after 20 generations. From Fig. 6, we can observe that our multi-objective genetic algorithm can find many solutions on the concave Pareto front.

We also applied a single-objective genetic algorithm where the weights \( w_1 \) and \( w_2 \) were fixed as follows:

\[ w_1 : w_2 = 100 : 1, 50 : 1, 20 : 1, 15 : 1, 10 : 1, 5 : 1, 2 : 1, \\
1 : 1, 1 : 2, 1 : 5, 1 : 10, 1 : 15, 1 : 20, 1 : 50, 1 : 100.\]
The single-objective genetic algorithm with these 15 different weight values found only two Pareto solutions: \((f_1, f_2) = (2, 4)\) and \((f_1, f_2) = (4, 1)\). From these simulation results, we can see that the single-objective genetic algorithm with constant weights cannot find the concave Pareto front even if various weight values were employed.

4. SIMULATION RESULTS FOR FLOWSHOP SCHEDULING

In this section, we demonstrate the effectiveness of our multi-objective genetic algorithm by computer simulations on a flowshop scheduling problem with two objectives: to minimize the makespan and to minimize the total tardiness. We also apply our algorithm to the flowshop scheduling problem with three objectives: to minimize the makespan, to minimize the total tardiness, and to minimize the total flowtime.

Flowshop scheduling problems are one of the most well-known problems in the area of scheduling. General assumptions of the flowshop scheduling problems can be written as follows (see Dudek et al. [16]). Jobs are to be processed on multiple stages sequentially. There is one machine on each stage. Machines are available continuously. A job is processed on one machine at a time without preemption, and a machine processes no more than one job at a time. In this paper, we assume that \(n\) jobs are processed in the same order on \(m\) machines. This means that our flowshop scheduling is the \(n\)-job sequencing problem. In this paper, the sequence of the \(n\) jobs is denoted by a string \(x = (x_1, \ldots, x_k, \ldots, x_n)\) where \(x_k\) is the \(k\)-th job to be processed on the \(m\) machines.

4.1. Two-objective flowshop scheduling problem

In computer simulations, we employed two objectives as scheduling criteria; to minimize the makespan (i.e., the completion time of the last job) and to minimize the total tardiness (i.e., the sum of the tardiness for the duedate of each job). We specified the due date of each job by the following procedure:

Step 1: Randomly generate a permutation of the \(n\) jobs.
Step 2: Calculate the completion time \(C_j\) of each job, \(j = 1, 2, \ldots, n\).
Step 3: Add a random integer \(\text{rnd}_j\) in the closed interval \([-100, 100]\) to each \(C_j\). That is, the duedate \(d_j\) of the \(j\)-th job is \(d_j = C_j + \text{rnd}_j\).

It is known that there is no correlation between the two objectives: the makespan and the total tardiness. In our multi-objective genetic algorithm, the fitness function \(f(x)\) can be written as

\[
  f(x) = -W_{\text{makespan}} \cdot \text{Makespan}(x) - W_{\text{tardiness}} \cdot \text{Tardiness}(x),
\]

where \(W_{\text{makespan}}\) and \(W_{\text{tardiness}}\) are the weights for the makespan and the total tardiness, respectively.

![Fig. 7. Comparison of our multi-objective genetic algorithm with two trials of the single-objective genetic algorithm.](image)
where \( \text{Makespan}(x) \) is the makespan when the \( n \) jobs are processed in the order of \( x \), \( \text{Tardiness}(x) \) is the total tardiness, and \( W_{\text{Makespan}} \) and \( W_{\text{Tardiness}} \) are non-negative variable weights for \( \text{Makespan}(x) \) and \( \text{Tardiness}(x) \), respectively. In computer simulations, \( W_{\text{Makespan}} \) and \( W_{\text{Tardiness}} \) were specified in the same manner as (3). Because \( \text{Makespan}(x) \) and \( \text{Tardiness}(x) \) should be minimized, the negative sign "−" is attached to each weight in (12).

As a test problem, we generated a flowshop scheduling problem with 20 jobs and 10 machines by randomly specifying the processing time of each job at each machine as an integer in the closed interval \([1, 99]\). In computer simulation, we used the following parameter specifications.

Population size: \( N_{\text{pop}} = 10 \),
Crossover probability: \( P_c = 1.0 \),
Mutation probability: \( P_m = 1.0/\text{string} \).

Non-dominated solutions obtained by the proposed algorithm are shown by \( \circ \) in Fig. 7 where the horizontal and vertical axes are the makespan and the total tardiness, respectively. In Fig. 7, non-dominated solutions obtained by the two trials of a single-objective genetic algorithm are shown by ■ (obtained by the genetic algorithm for minimizing the makespan) and ▲ (obtained by the genetic algorithm for minimizing the total tardiness). In the single-objective genetic algorithm, the fitness function was equivalent to the value of its objective function with the negative sign "−". A tentative set of Pareto optimal solutions was stored and updated in the single-objective genetic algorithm in the same manner as in the multi-objective genetic algorithm. In order to compare the proposed algorithm with the two trials of the single-objective genetic algorithm under the same computation load, we specified the number of evaluations of the fitness function as 100,000 in our multi-objective genetic algorithm and as 50,000 in each trial of the single-objective genetic algorithm. Therefore 100,000 solutions were evaluated in each approach.

From Fig. 7, we can see that the set of the non-dominated solutions obtained by our multi-objective genetic algorithm (\( \circ \)) is superior to the set of the non-dominated solutions obtained by the single-objective genetic algorithm (■ and ▲). This is because many solutions denoted by ■ and ▲ are dominated by solutions denoted by \( \circ \). This demonstrates the high performance of our multi-objective genetic algorithm.

The effectiveness of the elite preserve strategy is demonstrated in Fig. 8. In Fig. 8, "no elite", "2 elite" and "3 elite" mean that no elite individual, two elite individuals and three elite individuals
Multi-objective genetic algorithm

![Graphs showing comparison between multi-objective genetic algorithm, Schaffer's VEGA, and single-objective genetic algorithm.](image)

(a) The multi-objective genetic algorithm

(b) Schaffer’s VEGA

(c) The single-objective genetic algorithm

Fig. 9. Comparison of our multi-objective genetic algorithm with Schaffer’s VEGA and the single-objective genetic algorithm.

were preserved to the next population in the multi-objective genetic algorithm, respectively. In the "2 elite" algorithm, only the elite individuals with respect to the two objective functions were preserved. On the other hand, an individual randomly selected from a tentative set of Pareto optimal solutions was preserved in addition to such two elite individuals in the "3 elite" algorithm.
Table 1. Average CPU time used by each algorithm for finding the Pareto-optimal solutions in Fig. 9

<table>
<thead>
<tr>
<th>Algorithm</th>
<th>Proposed algorithm</th>
<th>Schaffer's VEGA</th>
<th>SOGA</th>
</tr>
</thead>
<tbody>
<tr>
<td>CPU time (s)</td>
<td>3.08</td>
<td>3.00</td>
<td>2.48</td>
</tr>
</tbody>
</table>

From Fig. 8, we can see that the "3 elite" method found better solutions than the "2 elite" and "no elite" methods. This means that our elite preserve strategy had an effect on the performance of the multi-objective genetic algorithm.

We also applied Schaffer's VEGA [11] and another version of the single-objective genetic algorithm to the same flowshop scheduling problem. In the single-objective genetic algorithm, we used the weights $W_{\text{Makespan}} = 5$ and $W_{\text{Tardiness}} = 2$ to calculate the fitness value. A tentative set of Pareto optimal solutions was stored and updated in the single-objective genetic algorithm in the

![Fig. 10. Comparison of our multi-objective genetic algorithm with Schaffer's VEGA and the single-objective genetic algorithm.](image-url)
same manner as in the multi-objective genetic algorithm. As a stopping condition, we used the total number of evaluations of strings (i.e., solutions). When 100,000 solutions were evaluated in each algorithm, the algorithm was terminated. The simulation results by the proposed genetic algorithm, Schaffer's VEGA and the single-objective genetic algorithm are shown in Figs 9(a), (b) and (c), respectively. We applied each algorithm five times to the same flowshop scheduling problem. Each algorithm began to search Pareto-optimal solutions from the same initial generation. From Fig. 9, we can see that better solutions were obtained by the multi-objective genetic algorithm because many solutions obtained by Schaffer's VEGA in Fig. 9(b) are dominated by multi-objective genetic algorithm solutions in Fig. 9(a). The single-objective genetic algorithm could find some better solutions than our multi-objective genetic algorithm, but the single-objective algorithm tended to fail to find large Pareto fronts. The average CPU time used by each algorithm for finding the solutions in Fig. 9 is shown in Table 1. We can see from Table 1 that a similar computation time was spent by each algorithm in computer simulations.

4.2. Three-objective flowshop scheduling problem

We also applied our multi-objective genetic algorithm to a flowshop scheduling problem with three objectives: to minimize the makespan, to minimize the total tardiness and to minimize the total flowtime (i.e., the sum of completion time over all jobs). We used the same parameters as used for the two-objective flowshop scheduling problem. Because it is difficult to show solutions in the three-dimensional objective space, we show the solutions by projecting them onto two-dimensional objective spaces. Figure 10 shows the simulation results obtained by our multi-objective genetic algorithm, Schaffer's VEGA and the single-objective genetic algorithm. In the single-objective genetic algorithm, we used the constant weights \( W_{\text{Makespan}} = 5, W_{\text{Tardiness}} = 2 \) and \( W_{\text{Flowtime}} = 1 \) to calculate the fitness value. From Fig. 10, we can observe that our multi-objective genetic algorithm could find a better set of solutions.

5. CONCLUSION

In this paper, we proposed a framework of genetic algorithms for multi-objective optimization problems. Our approach has two characteristic features. One is that the weights used for combining multiple objectives into a scalar fitness function are randomly specified for each selection. That is, the weights are not constant but variable in our multi-objective genetic algorithm. The other characteristic feature is that multiple elite individuals selected from a tentative set of Pareto optimal solutions are inherited to the next generation. By computer simulations, we showed that our multi-objective genetic algorithm could find concave Pareto optimal solutions, and we demonstrated that our multi-objective genetic algorithm could find better solutions than the VEGA (Vector Evaluated Genetic Algorithm) by Schaffer [11] and the single-objective genetic algorithm.

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