Evolutionary Algorithms with Segment-Based Search for Multiobjective Optimization Problems

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Abstract—This paper proposes a variation operator, called segment-based search (SBS), to improve the performance of evolutionary algorithms on continuous multiobjective optimization problems. SBS divides the search space into many small segments according to the evolutionary information feedback from the set of current optimal solutions. Two operations, micro-jumping and macro-jumping, are implemented upon these segments in order to guide an efficient information exchange among “good” individuals. Moreover, the running of SBS is adaptive according to the current evolutionary status. SBS is activated only when the population evolves slowly, depending on general genetic operators (e.g., mutation and crossover). A comprehensive set of 36 test problems is employed for experimental verification. The influence of two algorithm settings (i.e., the dimensionality and boundary relaxation strategy) and two probability parameters in SBS (i.e., the SBS rate and micro-jumping proportion) are investigated in detail. Moreover, an empirical comparative study with three representative variation operators is carried out. Experimental results show that the incorporation of SBS into the optimization process can improve the performance of evolutionary algorithms for multiobjective optimization problems.

Index Terms—Multiobjective optimization, hybrid evolutionary algorithms, variation operators, segment-based search.

I. INTRODUCTION

Many real-world problems involve simultaneous optimization of several competing criteria or objectives: Often, there is no single optimal solution, but rather a set of alternative solutions. Evolutionary algorithms (EAs) have been recognized to be suitable to deal with these multiobjective optimization problems (MOPs) due to their population-based property of achieving an approximation of the Pareto set in optimization problems. It is well-known that pure EAs may not be well suited for some complex problems [16]. The hybridization of EAs with other search techniques (e.g., variation operators, optimization algorithms, and machine learning methods) can greatly improve the search ability [32], [48], [60]. Aiming to obtain a balance between exploration and exploitation, hybrid EAs represent one of the growing research areas in the evolutionary computation community. Recent studies on hybrid EAs have demonstrated their usefulness on various problem domains [17], [41], [59].

Incorporating new variation operators into EAs is one important research topic in the area of hybrid EAs. According to the property of converging into a local optimum, variation operators can be divided into two categories: local search (LS) operators and non-LS operators. LS, which tries to guide solutions to converge into a local optimum, is a metaheuristic method. In general, an LS method starts from a certain point in the search space and moves it iteratively in a descent direction until a local optimum of the function is reached [27]. Hybridization of EAs with LS is known as memetic algorithms (MAs) [5], [28], [49]. Combining probabilistic search and deterministic search, MAs have been proven to be successful in solving complex optimization problems [14], [31], [37], [55].

One important application area of MAs is in the EMO community [15], [25]. Since the mid-1990s, a number of multiobjective MAs (MOMAs) have been proposed [11], [23], [45], [50]. For example, Ishibuchi and Murata [21] proposed a multiobjective genetic LS approach for flow shop scheduling, which is the first well-known MOMA. Jaszkiewicz [22] developed a weighted sum function-based MOMA, and suggested that scalarizing functions are particularly better at encouraging diversity than dominance ranking methods. The memetic-based Pareto archived evolution strategy (M-PAES), presented by Knowles and Corne [26], is an MOMA based on grid techniques. Unlike the aforementioned two algorithms, M-PAES does not use scalarizing functions but rather a Pareto ranking-based selection. Recently, Bosman [3] presented an analytical description of the set of all nondominated improving directions for any point in the decision space of an MOP. Moreover, he used this description in a gradient-based optimization algorithm.

On the other hand, some interesting extensions of established EMO algorithms with respect to the idea of MAs have been emerging. These extensions focus on some popular algorithms, such as the nondominated sorting GA II (NSGA-II) [6], strength Pareto EA 2 (SPEA2) [64], decomposition-based multiobjective EA (MOEA/D) [63], S metric selection EMO algorithm (SMS-EMOA) [2], and multiobjective covariance matrix adaptation evolution strategy (MO-CMA-ES) [20]. By integrating efficient LS strategies, these extensions greatly enhance the search ability of algorithms in comparison with their original versions (see [27], [36], [47], [56], [58]). Among these, the convergence acceleration operator (CAO) [1] and hill climber with sidestep (HCS) [30] are two representative local searchers. The former, based on a neural network technique, is
used to accelerate the convergence speed of EMO algorithms by mapping the improved vectors of the objective space into the decision space. The latter utilizes the geometry of the directional cones of optimization problems and works with or without the gradient information.

Hybridization of EAs with non-LS operators has also attracted increasing attention over the past few years. Despite the lack of the property of converging into a local optimum, these variation operators can clearly enhance the search ability of an EA when an appropriate combination between them and original genetic operators (such as crossover and mutation) in the EA is made. By exploiting specialized knowledge of the problem being solved, such hybrid EAs have been found to be promising in dealing with various optimization problems. Specifically, Tan et al. [51] proposed a two-phase hybrid evolutionary classification technique to extract classification rules in clinical practice for better understanding and prevention of unwanted medical events. Fan and Lampinen [12] incorporated differential evolution with a trigonometric mutation operator. Due to the capability of obtaining an acceptable solution with a lower number of objective function evaluations, this hybrid algorithm can be used in some expensive real-world problems, e.g., neural networks [12]. In order to address constrained optimization problems, Wang et al. [57] developed a hybrid EA which uses the idea of multiobjective optimization to balance exploration and exploitation during the evolutionary process. Tizhoosh [53] proposed an opposition-based learning (OBL) operator for intelligent optimization algorithms. Incorporated into differential evolution, particle swarm optimization, and estimation of distribution algorithms, OBL has been found to be competitive in many fields [42], [43].

For multiobjective optimization, it is also not uncommon to apply non-LS operators to improve the performance of EMO algorithms. Nebro et al. [40] presented a hybrid EMO algorithm that incorporates traditional mutation and crossover into the scatter search technique. In order to adapt the dynamic search of evolutionary operation in multiobjective optimization, Tan et al. [52] developed an adaptive variation operator that exploits the chromosomal structure of binary representation and synergizes the function of crossover and mutation. Elhossini et al. [10] hybridized PSO with genetic operators to solve MOPs. Genetic operators are modified to preserve the position information required by the PSO algorithm. Chan et al. proposed a hybrid EMO algorithm which uses the “jumping gene” operator to enhance the search ability during the evolutionary process for both binary-valued and real-valued scenarios [4], [39].

Overall, the impressive record of hybrid EAs with both LS and non-LS operators has proven their merits in the EMO community. Nevertheless, further study on hybrid EMO algorithms is still greatly needed to deeply explore and exploit the characteristics of multiobjective optimization. This paper focuses on the hybridization of EMO algorithms with non-LS operators and tries to develop a variation operator to improve the performance of EMO algorithms on continuous MOPs by utilizing the characteristics of multiobjective optimization.

A conspicuous feature of multiobjective optimizers is to consider the diversity of solutions in the evolutionary process. In this regard, performing single solution-based search as in single-objective optimization may fail if the information of other solutions is ignored. In fact, the essential characteristic in multiobjective optimization is that there exists no single optimal solution but a set of nondominated solutions due to the conflicting nature of multiple criteria. Therefore, in general, during the evolutionary process of a population, there is no solution better than all other solutions for all objectives, i.e., a solution may be optimal with respect to some objectives but not to the others. In this case, a variation strategy based on the search among “good” solutions may be more beneficial in multiobjective optimization because their offspring may inherit distinct strengths from them.

In this paper, we propose a variation operator, called segment-based search (SBS). SBS divides the search space into many small segments according to the evolutionary information feedback from the current optimal solutions (i.e., nondominated solutions), and performs information exchange among “good” individuals, which is called macro-jumping operation, and the local exploitation of a “good” individual, which is called micro-jumping operation, inside these segments. This is done in order to offer, on the one hand, an extensive probe in the space achieved already by the current evolutionary population, and on the other hand, a local refinement in the neighborhood of a single individual. In addition, the running of SBS is adaptive according to the evolutionary status of a population. It is activated only when general genetic operators of an algorithm do not work well, thus providing a needed fuel to the stagnant population without hindering the fast-evolving process of the population.

The remainder of the paper is organized as follows. The proposed operator, including the fundamental mechanisms of SBS and specific implementation of SBS in an EMO algorithm, is presented in Section II. Section III gives the algorithm setting, test functions, and performance metrics used for the experimental study in this paper. The performance validation of SBS and the comparison with other three operators are shown in Section IV and Section V, respectively. Finally, Section VI provides some concluding remarks along with pertinent observations and discussions for future work.

II. THE PROPOSED METHOD

It is well-established that the key design point for the hybridization of EAs with variation operators lies in the successful promotion of competition and cooperation between the forces of evolution and variation of individuals. When the balance between general genetic operators and introduced variation operators is not appropriately made, the performance of hybrid EAs is often severely degraded. This balance naturally involves several important issues, which should be addressed by the practitioner and are described as follows:

- What variation operator should be used in an EA for a particular problem?
- How often should the newly-introduced variation be applied (i.e., the frequency of variation operations)?
- Which individuals should be selected to perform the variation?
A. Segment-Based Search (SBS)

Many existing variation operators act on a single individual and generate offspring by exploiting its neighborhood. However, with an MOP, there exist some individuals that perform well for different objectives. Therefore, search based on information exchange among “good” individuals seems to be beneficial to producing better offspring. Here, we concentrate on the whole nondominated solution set in the current population. To be precise, we consider the region determined by the boundaries of the nondominated set in each dimension of the decision space, and divide it into a certain number of segments. Consequently, the SBS operator, specifically divided into micro-jumping and macro-jumping operations, will be implemented on the basis of these segments. Some segment-related concepts are given as follows.

For the convenience of discussion, we take a $D = 1$ dimensional decision space as an example. Figure 1 illustrates the configuration of segments in it.

First, the minimum and maximum decision values among a set of points $P$ are found and denoted as $\min(P)$ and $\max(P)$, respectively. The length of each segment will then be determined according to the following formula:

$$|S| = (\max(P) - \min(P))/L$$  \hspace{1cm} (1)

where $L$ is an integer value and usually set by the size of the population. Then, the lower boundary $lb$ and upper boundary $ub$ of the region are respectively defined as follows:

$$lb = \min(P) - |S|$$  \hspace{1cm} (2)

$$ub = \max(P) + |S|$$  \hspace{1cm} (3)

Clearly, the region is divided into $(L + 2)$ equal segments. They are $S_0, S_1, ..., S_L$, and $S_{L+1}$. The serial number of the segment where a point, denoted as $p$, is located can be calculated as follows:

$$sn(p) = \lfloor (p - lb)/|S| \rfloor$$  \hspace{1cm} (4)

For example, in Fig. 1, the serial number of $p$ is 2.

Next, we detail the SBS variation, namely, the micro-jumping and macro-jumping operations, according to the above preliminary concepts. Micro-jumping is a local refinement scheme which searches in the neighborhood of a solution, while macro-jumping is an extensive probe which explores in the whole specified region. Specifically, for a point $p$, a micro-jumping operation is to generate a random point in the range $[p - |S|, p + |S|]$; while for a macro-jumping operation, $p$ is transferred to one of the $L + 2$ segments in a random manner. That is, an integer $i$ among $0, 1, ..., L, L + 1$ is firstly randomly selected, and then a new point (denoted as $p'$) is formed according to the following formula:

$$p' = p + (i - sn(p)) \times |S|$$  \hspace{1cm} (5)

Clearly, macro-jumping is a jump to a point chosen at random from a limited set of points distributed across the whole search region. For example, in Fig. 1, assuming that $i$ is 5, $p$ will jump rightwards 3 segments to $p'$.

Note that the SBS operator extends the boundaries of the set of points: the new points generated by SBS have the opportunity of being located in segment $S_0$ or segment $S_{L+1}$. This slight relaxation of the interested region will be beneficial to enhancing the diversity of individuals in the evolution search, especially for multiobjective optimization. We will give an explanation regarding this point in the next section.

B. SBS in Multiobjective Optimization

In hybrid EAs, newly-introduced variation operation inevitably competes with traditional genetic operation for limited computational resources. Given a fixed computational budget, the frequency of new variation operation will directly affect the degree of traditional genetic operation. Clearly, more intense individual variation provides a greater chance of discovering promising solutions but limits the amount of evolution without incurring excessive computational resources. Therefore, care should be taken in setting the frequency of variation operations to utilize limited computational resources for achieving the best search performance.

In multiobjective optimization, comparing individuals is often based on their Pareto dominance relation. Given two individuals $a$ and $b$, $a$ is said to dominate $b$ if and only if $a$ is at least as good as $b$ in all objectives, and better in at least one objective. Accordingly, individuals in a population that are not dominated by any other individuals are denoted as the nondominated individuals with respect to this population.

For a genetic operator, if there exist newly generated individuals dominating some individuals of their parent population, it means that the operator has the capability of improving the performance of the current evolutionary population. And if there is no individual in the new population that dominates any individuals in their parent population, it may mean that the operator currently fails to provide enough power to drive the population forwards. Naturally, the number of nondominated individuals in the mixed set consisting of a population and its offspring population will largely reflect the evolutionary status of an EMO algorithm. At the initial stage of evolution,
it is usually easy to produce new individuals by the genetic operators (such as crossover and mutation operators) that dominate some individuals in their parent population, and thus, the number of nondominated individuals in the mixed population is often small. With the progress of evolution, that number will be gradually increased since there will be more and more nondominated individuals in the parent population, and it becomes harder for new individuals to dominate them. Especially, when the number of nondominated individuals exceeds a significant proportion of the mixed population, the convergence may slow down or even cease. This phenomenon often happens at the time the population approaches the (local) optimal front. This, to some extent, indicates that the current population is difficult to evolve further towards the desired area. Figure 2 gives two examples of the relationship between convergence and the number of nondominated individuals in the mixed set consisting of the parent and offspring populations during the evolutionary process of a typical run of the NSGA-II algorithm.

Clearly, when the number of nondominated individuals exceeds half of the size of the mixed set, the ability to evolve forwards will slow down to some extent. This occurs, for ZDT1 [65], when the population approaches the optimal front, whereas, for WFG1 [19], when the population might be trapped into some local optima in view of its relatively large GD value. In this study, we introduce SBS at this stage in order to enhance the exploration ability for the case of being trapped into local optima (e.g., for WFG1), or to refine the exploitation result for the case of approaching global optima (e.g., for ZDT1). The framework of integrating SBS into a general EMO algorithm is given in Algorithm 1.

In Algorithm 1, Step 4 tests the running condition of SBS. If the number of nondominated individuals in $R$ is larger than the population size $N$ ($|R| = 2N$), the SBS variation will be activated (Step 5); otherwise, $T$ becomes empty (Step 7). Step 9 implements the mating selection and genetic variation (e.g., crossover and mutation) of the original EMO algorithm to generate $(N - |T|)$ individuals. Therefore, a new population will be composed of $(N - |T|)$ individuals generated by genetic operators, and $|T|$ individuals generated by SBS (Step 10). Figure 3 graphically demonstrates the process of the case where SBS is activated.

The SBS variation only acts on the set $Q \setminus P$ (i.e., the nondominated individuals that have been eliminated by the environmental selection) rather than either $P$ (i.e., the elite individuals) or $R \setminus Q$ (i.e., the dominated individuals). There are $|Q \setminus P|$ individuals to be generated, but not all of them will be preserved. A new individual will enter the next population only if it is not dominated by its parent. Therefore, the number of evaluations per generation becomes higher when SBS is implemented (i.e., $|Q \setminus P| - |T|$ evaluations are added per generation).

Several strengths of applying SBS to the set $Q \setminus P$ are drawn as follows.

- Individuals from the set $Q \setminus P$ perform well in terms of performance.

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**Algorithm 1 EMO Algorithm with SBS**

**Require:** $N$ (population size)

1. Generate an initial population $P$ and create empty temporary sets $R$ and $P'$.
2. Create an empty temporary set $Q$ for storing the nondominated individuals in $R$, and an empty set $T$ for storing the individuals generated by SBS.
3. while termination criterion not fulfilled do
   4. if $|Q| > N$ then
      5. $T \leftarrow SBS\_variation(Q \setminus P, Q)$ /* Implement SBS when the size of $Q$ is larger than $N$ */
   6. else
      7. $T \leftarrow \emptyset$
   8. end if
   9. $P' \leftarrow Matings\_select\_variation(P, (N - |T|))$ /* Implement mating selection and genetic variation to generate $(N - |T|)$ individuals */
10. $P' \leftarrow P' \cup T$
11. $R \leftarrow P \cup P'$
12. $Q \leftarrow Nondominated\_select(R)$ /* Find all nondominated individuals of $R$ */
13. $P \leftarrow Environmental\_select(R)$
14. end while
15. return $P$

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1The results are evaluated by the generational distance (GD) metric [7]. GD assesses the convergence of a solution set by calculating the average Euclidean distance from the set to the Pareto front.

2The parameter settings of the algorithm are the same as in the experimental study, shown in Table II.

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**Fig. 2.** Evolutionary trajectory of the convergence metric GD for a typical run of NSGA-II on ZDT1 [65] and WFG1 [19]. The dashed line indicates the stages where the number of nondominated solutions (denoted as $|NS|$) in the mixed set $R$ (i.e., the combination of the parent and offspring populations in each generation) is smaller than or equal to the population size $N$ ($|R| = 2N$), and the solid line indicates the stages where $|NS|$ is larger than $N$. 

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3Note that the segments in the variation are defined using $Q$, cf. Step 3 in Algorithm 2.
of convergence and there are no individuals dominating them in the set \(R\). Applying SBS on them has a greater chance to generate better offspring in comparison with applying SBS on the dominated individuals in \(R \setminus Q\).

- Although being nondominated solutions in \(R\), individuals from \(Q \setminus P\) are not used as parents to generate offspring. Moreover, in contrast to the probable long time existence of the elite individuals in \(P\) during the evolutionary process, solutions in \(Q \setminus P\) will not exist in the next generation population. Therefore, applying SBS on them means to give them a chance to contribute their value, which may be beneficial to exploring the potential of every nondominated individual in \(R\).

- The number of individuals in \(Q \setminus P\) varies adaptively according to the evolutionary situation. On the one hand, when the size of the nondominated set \(Q\) is relatively small, the evolutionary population appears to still have some potential to be exploited and thus the impact of SBS is relatively weak. On the other hand, when the size becomes larger, the population in general evolves more slowly and the impact of SBS increases correspondingly.

Up to now, the four issues with respect to the design of a hybrid EA mentioned at the beginning of this section have been addressed—a segment-based variation operator is used; it is activated when the number of nondominated individuals exceeds half of the size of the mixed set consisting of the parent and offspring populations; it acts on nondominated individuals that are eliminated by the environmental selection; and it occurs before mating selection and genetic operations but after the environmental selection process. Algorithm 2 gives the pseudocode of the implementation of SBS.

The main procedure of SBS can be divided into two parts. The first part, consisting of Steps 2–5, makes preparations for the SBS variation, including setting up the environment of segments (i.e., the length of a segment and the lower and upper boundaries) in each dimension of the decision space and selecting individuals for variation. The second part, consisting of Steps 6–29, details the implementation process of SBS. In this part, there are two user-defined probability parameters, \(MiP\) and \(SR\). \(MiP\) denotes a proportion of micro-jumping versus macro-jumping, and \(SR\) denotes a rate of the SBS variation. Additionally, parameter \(jrand\) is a random integer to ensure that at least one element is selected to vary in each interested individual. Steps 26–28 are used to determine the survival of new individuals. A new individual will be saved only if it does not perform worse than its parent, i.e., it is not dominated by its parent.

Finally, it is worthwhile to mention that the range of the concerned region in SBS always changes in the evolutionary process since it is determined by the current population. This, similar to the OBL operator developed in [43], [53], will help to shrink the search space and employ existing knowledge of evolution. Nevertheless, a key difference from the OBL variation is the slight relaxation of the boundaries of the solution set in SBS. This will be more useful to enhance

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**Algorithm 2 SBS_variation**\((Q \setminus P, P)\)

**Require:** \(N\) (population size), \(D\) (problem dimension in the decision space), \(SR\) (SBS rate), \(MiP\) (micro-jumping proportion), \(MiM\) (micro-jumping mark), \(MaN\) (macro-jumping number), \(jrand\) (random parameter to ensure variation)

1. Create an empty temporary set \(U\) and an empty set \(T\) for storing the individuals generated by SBS
2. for all dimension \(j\) in the decision space, \(j = 1, \ldots, D\)
3. \(Set\_segment(Q, j, N)\)
4. end for
5. \(U \leftarrow Q \setminus P\)
6. for all \(p_i \in U, i = 1, \ldots, |U|\)
7. if \(jrand(0, 1) < MiP\)
8. then
9. \(MiM \leftarrow 1\)
10. else
11. \(MaN \leftarrow \lceil rand(0, 1) \times (N + 2) \rceil\)
12. end if
13. \(jrand \leftarrow \lceil rand(0, 1) \times D \rceil + 1\)
14. for all dimension \(j\) of \(p_i, j = 1, \ldots, D\)
15. if \(rand(0, 1) < SR \lor j = jrand\)
16. then
17. \(MiM = 1\)
18. \(p_{ij} \leftarrow rand[p_{ij} - |S|_j, p_{ij} + |S|_j]\)
19. else
20. \(p_{ij} \leftarrow p_{ij} + (MaN - Sn_j(p_{ij})) \times |S|_j\)
21. end if
22. \(p_{ij} \leftarrow p_{ij}\)
23. end if
24. end for
25. \(eval(\hat{p}_j)\)
26. if \(\neg (p_i < \hat{p}_j)\)
27. then
28. \(T \leftarrow T \cup \{\hat{p}_j\}\)
29. end if
30. end for
31. return \(T\)
the diversity of the population in multiobjective optimization because the optimal result of an MOP is a region, rather than a solitary point, in contrast to a single-objective optimization problem. An experimental study of this relaxation strategy will be given in Section IV-C.

III. EXPERIMENTAL DESIGN

In order to verify SBS, we integrate it into three widely-used EMO algorithms, NSGA-II [6], SPEA2 [64] and SMS-EMOA [2]. NSGA-II is one of the most popular EMO algorithms. The main characteristic is its fast nondominated sorting and crowding distance-based density estimation. SPEA2 is also a prevalent EMO algorithm, which borrows a so-called fitness strength value and the k-th nearest neighbor density estimation to select individuals into the next population. SMS-EMOA is an indicator-based EMO algorithm, which maximizes the hypervolume contribution of a population during the evolutionary process. Combined with the concept of nondominated sorting, SMS-EMOA can produce a well-converged and well-distributed solution set.

Note that the original SMS-EMOA adopts a steady-state evolution model, which creates only one new member and then tests whether it can enter the archive or not at each step of the algorithm. Here, to make SMS-EMOA suitable in the framework of SBS, we modify the original SMS-EMOA and adopt a generational evolution version of the algorithm, which, like NSGA-II and SPEA2, creates a set of new solutions and tests whether they can enter the next population or not at each step of the algorithm [8]. In addition, since the original implementation of SPEA2 and SMS-EMOA does not contain any constraint handling mechanism, we modified their implementation by applying the same constraint handling mechanism used in NSGA-II.

In the following subsections, we introduce the test problems, performance metrics, and parameter setting for the comparative study.

A. Test Problems

A comprehensive set of 36 MOPs are used for performance verification of the proposed method. These test problems have been commonly used in the literature. They can be divided into several groups according to the number of objectives and constraint conditions.

For the first group of bi-objective unconstrained problems, we first choose problems from Veldhuizen’s studies [54], including Schaffer, Fonsec1a1, Fonsec2a, Kursawe, and Poloni problems. Then, the ZDT problem family (including ZDT1, ZDT2, ZDT3, ZDT4, and ZDT6) [65] is considered. Finally, the walking fish group (WFG) problem family (WFG1 to WFG9) [19] based on variable linkage is also contained in this group. The second group consists of the bi-objective constrained problems from [7], including BEL, BINH, OSY2, TNK, SRIN, and Constr_Ex. For the third group, we select MOPs with three objectives. First, four Viennet problems [54] (VNT1, VNT2, VNT3, and VNT4) are chosen, of which the first three problems have three objectives and no constraints, and the fourth one has three objectives and three constraints.

Then, the unconstrained DTLZ problem family [9] (DTLZ1 to DTLZ7) is considered.

B. Performance Metrics

To compare the algorithms, we use two performance metrics, i.e., inverted generational distance (IGD) [63] and hypervolume (HV) [7], which are commonly used in the EMO community [34], [35]. In addition, median attainment surfaces [13] are provided for visual inspection of the performance of algorithms on some test problems. IGD measures the average distance from the points in the Pareto front to their closest solution in the obtained set. Mathematically, let $P^*$ be a reference set representing the Pareto front, then the IGD value from $P^*$ to the obtained solution set $P$ is defined as follows:

$$IGD = \sum_{z \in P^*} d(z, P) / |P^*|$$

where $|P^*|$ denotes the size of $P^*$ (i.e., the number of points in $P^*$) and $d(z, P)$ is the minimum Euclidean distance from point $z$ to $P$. A low IGD value is preferable, which indicates that the obtained solution set is close to the Pareto front as well as having a good distribution.

The HV metric calculates the volume of the objective space enclosed by the obtained solution set and a reference point. A large value is preferable. It can be described as the Lebesgue measure $\lambda$ of the union hypercubes $h_i$ defined by a solution $p_i$ in the set and the reference point $x_{ref}$ as follows:

$$HV = \lambda(\{ \bigcup_i h_i \mid p_i \in P \}) = \lambda( \bigcup_{p_i \in P} \{ x \mid p_i \prec x < x_{ref} \})$$

Note that, in the calculation of HV, the solutions that do not dominate the reference point are discarded (i.e., the solutions that are worse than the reference point in at least one objective will contribute zero to HV).

Plotting median attainment surfaces of EMO algorithms’ results (i.e., the final solutions) is a good performance comparison method, which allows visual inspection of the performance of algorithms regarding convergence and diversity [13]. For one run of an algorithm, the attainment surface of its results divides the objective space into two regions: one containing solutions which are dominated by its results, and another containing solutions that dominate its results. For multiple runs of an algorithm, the median attainment surface is the surface that the algorithm’s results are expected to attain in 50% of the runs. Here, we use Knowles’s approach to plot the median attainment surface [24]. The code is available at http://www.cs.man.ac.uk/~jknowles/plot_attainments/.

C. General Experimental Setting

The parameter settings for all conducted experiments are given as follows unless explicitly mentioned otherwise.

- **Parameter setting in SBS:** In the SBS operator, both SR and MiP (i.e., the SBS rate and micro-jumping proportion) are set to 0.5 for the comparative study. The effect of these two parameters on the performance of SBS will be investigated in detail in Section IV-D.
• Parameter setting of genetic operators in the tested algorithms: In NSGA-II, SPEA2, and SMS-EMOA, two widely-used crossover and mutation operators, i.e., the simulated binary crossover (SBX) and polynomial mutation [7], are chosen. Following the practice in [6], the distribution indexes in both the SBX and polynomial mutation are set to 20. A crossover probability $p_c = 1.0$ and a mutation probability $p_m = 1/D$ (where $D$ is the number of decision variables) are used according to [7].

• Population and archive size: Like most of the studies of EMO algorithms, the population size is set to 100, and the archive set, if used, is maintained with the same size as the population size.

• Number of decision variables in scalable problems: ZDT, DTLZ, and WFG are three suites of test problems, whose dimensionality $D$ can be specified by the user. The default dimensionality $D_{\text{default}}$ is set to 30 for the first three ZDT problems, to 10 for ZDT4 and ZDT6, and to 12 for all the DTLZ problems except DTLZ1 and DTLZ7, whose $D_{\text{default}}$ is set to 7 and 22, respectively. For the WFG suite, $D_{\text{default}}$ is set to 6.

• Number of runs and stopping condition: We independently run each algorithm 30 times for each test problem. The termination criterion of an algorithm is a predefined number of evaluations. Here, we set the number of evaluations to different values for problems with different numbers of objectives, since the difficulty of problems generally increases with the number of objectives [46]. The number of evaluations is set to 25,000 and 30,000 for bi-objective and tri-objective problems, respectively.

• Reference point setting in HV: Similar to [29], we select the integer point slightly larger than the worst value of each objective on the Pareto front of a problem as its reference point, as given in Table I.

• Substitution of the Pareto front for IGD: For the IGD metric, it is necessary to know the Pareto front of test problems. In most of the test problems used in this work, including families ZDT, DTLZ, and WFG, their Pareto fronts are known. For them, we select 10,000 evenly-distributed points along the Pareto front as its substitution in the calculation of IGD. For other test problems, the substitution of their Pareto fronts is available from the web site http://www.cs.cinvestav.mx/~emoobook/.

The parameter settings are summarized in Table II.

### Table I

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### Table II

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<th>Parameter settings in comparative experiments</th>
<th>S/H in SBS</th>
<th>Crossover Prob.</th>
<th>Population Size</th>
</tr>
</thead>
<tbody>
<tr>
<td>Crossover SBX</td>
<td>0.5</td>
<td>1.0</td>
<td>100</td>
</tr>
<tr>
<td>Mutation Polynomial</td>
<td>0.5</td>
<td>0.5</td>
<td>100</td>
</tr>
<tr>
<td>Archive size</td>
<td>20</td>
<td>Run Number</td>
<td>30</td>
</tr>
</tbody>
</table>

### D. Experimental Classification

In order to carry out a reliable and comprehensive comparison, we categorize the experiments into five groups:

1. Performance verification of SBS;
2. Effect of the problem dimensionality;
3. Effect of the boundary relaxation strategy;
4. Setting of newly added control parameters;
5. Comparison of SBS with OBL [43], [53], jumping gene [4], [39], and Mühlenbein’s mutation [38].

Unless mentioned otherwise, the same parameter settings as described in the previous subsection are adopted in each experiment.

### IV. Performance Verification of SBS

In this section, we investigate the performance of the proposed SBS operator following the experimental design outlined in the previous section. We embed SBS into NSGA-II, SPEA2, and SMS-EMOA, and call them NSGA-II+SBS, SPEA2+SBS, and SMS-EMOA+SBS, respectively. The values included in the tables of results are mean and standard deviation (SD) over 30 independent runs, and a better mean is highlighted in boldface. In addition, a t-test is used to compare tested algorithms on each function. The symbol “*” indicates that the $p$ value of 58 degrees of freedom is significant at a 0.05 level of significance by a two-tailed $t$-test.

### A. Performance Verification of SBS

First, we compare NSGA-II+SBS with the original NSGA-II. The results of 36 benchmark functions regarding the IGD and HV metrics are given in Table III and Table IV, respectively. Clearly, NSGA-II+SBS performs significantly better than NSGA-II regarding both metrics. NSGA-II+SBS obtains better values in 26 and 25 out of the 36 test problems regarding IGD and HV, respectively. Also, for the majority of the problems on which NSGA-II+SBS outperforms NSGA-II, the results have statistical significance (16 out of the 26 problems for IGD and 19 out of the 25 problems for HV). Note that the degree of performance improvement brought by SBS is considerable on some problems where NSGA-II fails to converge, such as WFG1 and DTLZ6. For these test cases, NSGA-II+SBS can greatly enhance the search ability, successfully guiding the solutions to approximate the Pareto front of the problem. This occurrence may be attributed to the macro-jumping operation in SBS which can produce some promising individuals to help the current stagnant population. Figure 4 plots the 50% attainment surfaces [24] across the 30 runs of NSGA-II and NSGA-II+SBS on DTLZ6. It is clear that NSGA-II+SBS performs better than NSGA-II since

The parameter settings are summarized in Table II.
the median attainment surface of NSGA-II+SBS has a larger dominated space than that of NSGA-II.

In order to investigate the “success rate” of the SBS operator during the evolutionary process, we also show, as a function of the number of evaluations, the probability that a solution generated by SBS has been inserted into the population across the 30 runs of NSGA-II+SBS on DTLZ6 in Fig. 5. Clearly, despite fluctuating, the probability value exceeds 30% most of the time. This means that SBS is an efficient operator to generate “good” solutions (i.e., solutions that are not dominated by their parent) during the evolutionary process.

To better understand the performance difference between NSGA-II and NSGA-II+SBS, we show the average HV result with standard errors during the whole evolutionary process across all 30 runs of the two algorithms on DTLZ6 and WFG1 in Fig. 6 and Fig. 7, respectively. As can be seen from the figures, an evident improvement of the HV value can be achieved when SBS is embedded into NSGA-II. For DTLZ6, the HV trajectory of NSGA-II+SBS increases rapidly from about 6,000 evaluations and approximates the optimal value (around 6.10) at about 23,000 evaluations. For WFG1, NSGA-II+SBS shows a clear advantage over NSGA-II from around 7,000 evaluations until the end.

Tables V and VI show the comparative results of SPEA2 and SPEA2+SBS. Similar to the case of NSGA-II+SBS, SPEA2+SBS has a clear advantage over the original SPEA2. It obtains a better value in 26 and 25 out of all 36 test
### TABLE V

<table>
<thead>
<tr>
<th>Problem</th>
<th>SPEA2</th>
<th>SPEA2+SBS</th>
<th>Problem</th>
<th>SPEA2</th>
<th>SPEA2+SBS</th>
<th>Problem</th>
<th>SPEA2</th>
<th>SPEA2+SBS</th>
</tr>
</thead>
</table>
| SCH     | 1.6602E–2 (1.1E–4)  | 1.7022E–2 (1.2E–4)
| FON1    | 4.6642E–3 (7.1E–5)
| FON2    | 4.1800E–3 (3.2E–5)
| FON3    | 5.1392E–3 (1.5E–4)
| ZDT1    | 4.1795E–3 (9.3E–5)
| ZDT2    | 4.1675E–3 (1.2E–4)
| ZDT3    | 5.5675E–3 (5.5E–3)
| ZDT4    | 6.5026E–3 (1.3E–4)
| ZDT6    | 8.3703E–3 (5.8E–4)
| WFG1    | 6.7301E–1 (1.5E–1)
| WFG2    | 1.2936E–2 (8.2E–3)

Indicates that the p-value of 58 degrees of freedom is significant at a 0.05 level of significance by a two-tailed t-test.

### TABLE VI

**HV Comparison (Mean and SD) of SPEA2 and SPEA2+SBS: A Better Mean for Each Case is Highlighted in Boldface**

<table>
<thead>
<tr>
<th>Problem</th>
<th>SPEA2</th>
<th>SPEA2+SBS</th>
<th>Problem</th>
<th>SPEA2</th>
<th>SPEA2+SBS</th>
<th>Problem</th>
<th>SPEA2</th>
<th>SPEA2+SBS</th>
</tr>
</thead>
</table>
| SCH     | 2.2275E+1 (6.9E–4)
| FON1    | 3.0621E+0 (1.4E–4)
| FON2    | 3.3368E+0 (1.4E–4)
| FON3    | 3.7071E+0 (1.1E–2)
| FON4    | 5.7319E+0 (1.8E–2)
| ZDT1    | 3.6596E+0 (4.4E–4)
| ZDT2    | 3.3244E+0 (1.7E–4)
| ZDT3    | 3.5266E+0 (3.4E–3)
| ZDT4    | 3.0192E+0 (1.0E–3)
| ZDT6    | 3.3024E+0 (3.4E–3)
| WFG1    | 7.6940E+0 (7.5E–1)
| WFG2    | 1.1099E+0 (4.1E–1)

Indicates that the p-value of 58 degrees of freedom is significant at a 0.05 level of significance by a two-tailed t-test.

### TABLE VII

**IGD Comparison (Mean and SD) of SMS-EMOA and SMS-EMOA+SBS: A Better Mean for Each Case is Highlighted in Boldface**

<table>
<thead>
<tr>
<th>Problem</th>
<th>SMS-EMOA</th>
<th>SMS-EMOA+SBS</th>
<th>Problem</th>
<th>SMS-EMOA</th>
<th>SMS-EMOA+SBS</th>
<th>Problem</th>
<th>SMS-EMOA</th>
<th>SMS-EMOA+SBS</th>
</tr>
</thead>
</table>
| SCH     | 1.9287E–2 (3.0E–4)
| FON1    | 6.3745E–3 (5.9E–5)
| FON2    | 4.2950E–3 (2.6E–4)
| FON3    | 3.3400E–2 (6.4E–4)
| FON4    | 5.3218E–2 (1.5E–3)
| ZDT1    | 3.7291E–3 (1.6E–5)
| ZDT2    | 4.5852E–3 (5.6E–5)
| ZDT3    | 4.3191E–3 (2.6E–4)
| ZDT4    | 4.8381E–3 (2.1E–4)
| ZDT6    | 4.9154E–3 (2.8E–4)
| WFG1    | 4.2020E+1 (2.1E–2)
| WFG2    | 1.1878E–2 (1.8E–3)

Indicates that the p-value of 58 degrees of freedom is significant at a 0.05 level of significance by a two-tailed t-test.

### TABLE VIII

**HV Comparison (Mean and SD) of SMS-EMOA and SMS-EMOA+SBS: A Better Mean for Each Case is Highlighted in Boldface**

<table>
<thead>
<tr>
<th>Problem</th>
<th>SMS-EMOA</th>
<th>SMS-EMOA+SBS</th>
<th>Problem</th>
<th>SMS-EMOA</th>
<th>SMS-EMOA+SBS</th>
<th>Problem</th>
<th>SMS-EMOA</th>
<th>SMS-EMOA+SBS</th>
</tr>
</thead>
</table>
| SCH     | 2.2281E+1 (2.0E–4)
| FON1    | 3.0631E+0 (5.1E–5)
| FON2    | 3.3374E+0 (2.7E–5)
| FON3    | 3.7131E+0 (1.3E–2)
| FON4    | 7.5412E–1 (5.7E–5)
| ZDT1    | 3.6616E+0 (6.0E–5)
| ZDT2    | 3.3292E+0 (5.2E–5)
| ZDT3    | 3.8151E+0 (9.1E–5)
| ZDT4    | 3.0305E+0 (3.0E–5)
| ZDT6    | 1.0114E+1 (1.0E+0)
| WFG1    | 1.1121E+1 (4.1E–1)

Indicates that the p-value of 58 degrees of freedom is significant at a 0.05 level of significance by a two-tailed t-test.
Fig. 6. Evolutionary trajectories of HV obtained by NSGA-II and NSGA-II+SBS on DTLZ6.

Fig. 7. Evolutionary trajectories of HV obtained by NSGA-II and NSGA-II+SBS on WFG1.

problems in terms of IGD and HV, respectively. Also, the difference of the two algorithms on most of the problems has statistical significance. Specifically, the proportion of the problems where SPEA2+SBS outperforms SPEA2 with statistical significance is 16/26 and 18/25 instances regarding IGD and HV, respectively. Additionally, the performance improvement is noticeable for some cases, e.g., WFG5 and WFG6. Figure 8 plots the 50% attainment surfaces across the 30 runs of SPEA2 and SPEA2+SBS on WFG6. Clearly, the median attainment surface of SPEA2+SBS has a larger dominated space than that of SPEA2. In fact, the median attainment surface of SPEA2+SBS already approximates the Pareto front of the problem, which means that the final solutions obtained by SPEA2+SBS can statistically achieve a good balance between convergence and diversity.

Tables VII and VIII show the comparative results between SMS-EMOA and SMS-EMOA+SBS. Although the advantage of SMS-EMOA+SBS over SMS-EMOA seems to be not as clear as that of NSGA-II+SBS over NSGA-II and of SPEA2+SBS over SPEA2, the proposed operator can improve SMS-EMOA for the majority of the problems. Specifically, SMS-EMOA+SBS outperforms SMS-EMOA in 21 and 22 out of the 36 test problems regarding IGD and HV, respectively. Also, for most of the problems on which SMS-EMOA+SBS performs better, the results have statistical significance (16 out of the 21 problems for IGD and 17 out of the 22 problems for HV). In contrast, the number of problems where SMS-EMOA outperforms SMS-EMOA+SBS with statistical significance is 8 and 9 in terms of IGD and HV, respectively.

It is important to note that the influence of SBS is different for different kinds of test problems. Integrated by the proposed operator, NSGA-II and SPEA2 achieve a clear improvement for most of the ZDT and DTLZ problems. For the WFG family, the proposed operator works well on some problems where the algorithm fails to approach the Pareto front, such as WFG1, WFG5, and WFG6, but performs poorly on some relatively easy problems, such as WFG3 and WFG4.

The UF and CF family [62] is also a set of popular test problems, which emphasize the complexity of the shape of the Pareto set. In some preliminary experiments, we observed that for the UF and CF problem family, the performance of the tested algorithms (i.e., NSGA-II, SPEA2, and SMS-EMOA) has not a clear improvement when integrated with the proposed SBS operator. This may mean that SBS is not well suited for MOPs with the Pareto set having such complex shape in the decision space.

**Result summary**—From the three groups of experiments on 36 benchmark functions, the algorithms with the integration of the SBS operator show a clear improvement. For most of the test problems, a better balance between convergence and diversity can be achieved when SBS is integrated into the algorithms.

B. Effect of Dimensionality

The experimental results in the previous section have shown that the SBS operator can improve the performance of EMO algorithms. Next, we study the influence of parameter setting on the performance of algorithms. Due to space limitation, we only consider the implementation of SBS into NSGA-II and present the HV results of NSGA-II and NSGA-II+SBS in this and the following sections. Similar results can also be observed for the other algorithms.

This section investigates the effect of problem dimensionality (i.e., the number of decision variables) on the performance of NSGA-II+SBS. The same experiments are repeated for $D = 2D_{defaul}$ on each scalable function from our test set. All other control parameters are kept unchanged. Table IX gives the results on all 21 scalable functions. Note that for some problems, the HV value of the obtained solution set may be equal to zero, which means that no solutions in the set can converge into the area determined by the reference point. In the case, the assessment result fails to indicate the performance difference between algorithms. To solve this issue, we reset the reference point for these problems (marked by * in the table) by specifying the integer point slightly larger than the
As can be seen from the table, NSGA-II+SBS outperforms NSGA-II in 16 out of the 21 problems, whereas only for WFG4 and DTLZ5, NSGA-II outperforms NSGA-II+SBS with statistical significance. On the other hand, it is clear that as the number of decision variables increases, both algorithms perform worse on all functions except on WFG8, where, interestingly, higher dimensionality seems to be more suitable. In addition, by a comparative observation in Tables IV and IX, it can be found that the advantage of NSGA-II over NSGA-II+SBS varies with the increase of problem dimensionality on some test problems. For problems WFG3 and DTLZ7 where NSGA-II outperforms NSGA-II+SBS under the default settings, NSGA-II+SBS achieves a better result when $D = 2D_{\text{default}}$.

In real-world scenarios, it is not uncommon to face a problem with hundreds of decision variables. To verify the efficiency of SBS for a problem with a high dimensionality, we compare the performance between the two algorithms on all problems with dimensions 100 and 1,000, as shown in Tables X and XI, respectively. Interestingly, from the tables, the advantage of NSGA-II+SBS over NSGA-II becomes clearer when a high number of decision variables are involved. For both $D = 100$ and $D = 1000$, NSGA-II+SBS performs better than NSGA-II in 18 out of the 21 problems. Also, for most of the problems on which NSGA-II+SBS outperforms its competitor, the results have statistical significance (14 out of the 18 problems when $D = 100$ and 15 out of the 18 problems when $D = 1000$).

In addition, the investigation of problem dimensionality in the objective space is also an important issue, which we leave for future study. In this context, we plan to apply SBS to some recent evolutionary many-objective optimization algorithms, such as GrEA [61] and SPEA2+SDE [33], and test the performance of the proposed operator in a high-dimensional objective space.

**Result summary**—While the algorithm integrated with the SBS operator obtains a lower HV value on most of the scalable functions with the increase of dimensionality, it still maintains stable advantage over its original version. Moreover, the advantage becomes clearer when the problem...
dimensionality increases, with NSGA-II+SBS outperforming NSGA-II on around 85% of all the problems when $D = 100$ and $D = 1000$.

C. Effect of the Boundary Relaxation Strategy

In this section, we investigate the effect of the boundary relaxation strategy (BRS) in SBS by 1) implementing SBS without BRS and 2) expanding the boundary in BRS. For this purpose, all parts of the proposed SBS operator are kept untouched except that the boundary is adjusted.

First, considering the SBS without BRS, Eqs. (2) and (3) will be changed into $lb = \min(P)$ and $ub = \max(P)$, respectively, and there will be $L$, instead of $(L+2)$, segments to be considered in each dimension ($L$ is set to $N$, i.e., the population size).

Table XII gives the HV values of NSGA-II+SBS without BRS on all the 36 test problems, and the results of NSGA-II+SBS from Table IV are also repeated in this table to ease the comparison between them. Clearly, the algorithm without BRS performs significantly worse than the original NSGA-II+SBS. The former outperforms the latter in 10 out of the 36 problems, and with statistical significance for only one problem (TNK). On the other hand, NSGA-II+SBS with BRS achieves a better HV value in 25 out of the 36 problems, and with statistical significance for more than half of these problems (15 of the 25 problems). In addition, on the problem FON1, both algorithms have an equal result regarding the mean and SD values.

Next, we test the effect of the boundary relaxation degree in SBS to the algorithm’s performance. Specifically, the upper and low boundaries are expanded to $lb = \min(P) - k|S|$ and $ub = \max(P) + k|S|$, respectively, where $k$ is set to 2, 3, 5, and 10. Therefore, there are $L + 2k$ segments to be considered in each dimension. Interestingly, we observe that the effect of the boundary relaxation degree is different for different problems. Here, we demonstrate the results of NSGA-II+SBS on three WFG test problems, WFG1, WFG2, and WFG8. Figure 9 plots the HV curve of the algorithm with different degrees of boundary relaxation in SBS. Clearly, the optimal setting of $k$ is varying, with $k = 5$, $k = 10$, and $k = 1$ for WFG1, WFG2, WFG8, respectively. This means that a good tradeoff between extensive exploration and local exploitation is distinct for different problems.

Result summary—After eliminating the BRS in SBS, the HV value of the algorithm drops on 25 problems, and has a slight improvement on only 10 problems. This clearly demonstrates that BRS is useful to enhance the search ability in multiobjective optimization. However, the optimal setting of the boundary relaxation degree is varying for different problems. Adaptive setting according to the evolutionary information may be a promising direction in the future study.

D. Effects of the SBS Rate (SR) and Micro-jumping Proportion (MiP)

In SBS, two probability parameters, $SR$ and $MiP$, are introduced to control the rate of the SBS variation and the proportion of micro-jumping versus macro-jumping, respectively. Although both parameters were fixed for all experiments, the performance of SBS can vary with their different settings. In this section, we investigate the effects of $SR$ and $MiP$, and try to provide a proper value (or range) of them for users. Here, a representative function set, the WFG problem family, is selected for demonstration.

To study the sensitivity of SBS to $SR$ and $MiP$, we repeat the experiments conducted in Section IV-A
for $SR, MiP \in [0, 1]$ using a step size of 0.1 (i.e., 30 trials per function for each combination of $SR, MiP \in \{0, 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1.0\}$). Due to space limitation, we do not show the results for all WFG problems. Instead, we show only the results for two representative functions WFG1 and WFG3 with respect to the 121 different combinations of $SR$ and $MiP$ in Fig. 10. Moreover, for comparison, we also show the HV value obtained by the original NSGA-II in the figure.

It is clear from Fig. 10 that the performance of NSGA-II+SBS varies drastically with $SR$ and $MiP$. For WFG1, NSGA-II+SBS works well when both $SR$ and $MiP$ are around 0.5, and interestingly, NSGA-II+SBS with any settings of the two parameters significantly outperforms the original NSGA-II (the worst HV value of NSGA-II+SBS in the 121 combinations is 8.7250, which is larger than the HV value 8.2253 of NSGA-II). For WFG3, a better HV can be obtained by the algorithm when working with a lower $SR$ or a higher $MiP$ value. The best and worst HV values of NSGA-II+SBS in the 121 combinations are 10.9376 and 10.9127, respectively. And the HV value for NSGA-II is 10.9333, which is higher than most of the results obtained by NSGA-II+SBS under different combinations of the parameters. In the following, we separately investigate the two parameters, trying to give a proper setting of them for users.

First, we consider the parameter $SR$. We repeat the experiments conducted in Section IV-A for $SR \in [0, 1]$ with a step size of 0.1. All other control parameters are kept unchanged. Figure 11(a) gives the results of NSGA-II+SBS with different settings of $SR$ on the WFG problem family, and the results of NSGA-II are also shown for comparison. Note that $SR = 0$ does not mean that SBS is not performed, but rather it means that only one dimension of the decision space is selected to vary (i.e., $j = \text{rand}$ from Step 15 in Algorithm 2). It is clear from the figure that the variation degree of performance has an apparent distinction on different problems with respect to different settings of the parameter. For some problems, such as WFG1, WFG2, and WFG8, the HV value varies with different settings of $SR$; but for some other problems, such as WFG3, WFG4, WFG7, and WFG9, the HV value appears to be relatively invariant. This occurrence is due to the difference of the performance of NSGA-II on the WFG problems. For WFG1, WFG2, and WFG8, the solutions obtained by NSGA-II are far from the Pareto front. A clear improvement of the HV result can be achieved when setting $SR$ and $MiP$ properly in SBS. For WFG3, WFG4, WFG7, and WFG9, the solutions of NSGA-II are already quite close to the Pareto front. So whatever setting for $SR$ and $MiP$, the degree of change of the HV result is relatively apparent, compared with that on WFG1, WFG2, and WFG8.

In order to investigate the effect of $SR$ on all the WFG problems more clearly, we show the HV value with finer scales for some problems where it is difficult to identify in Fig. 11(a). The results of WFG5 and WFG6 are given in Fig. 11(b). The results of WFG3, WFG4, WFG7, and WFG9 are plotted in Fig. 11(c), where the HV value of WFG3 and WFG9 is modified by adding or subtracting a constant to integrate them into one figure as follows: $HV_{WFG3} = HV_{WFG3} - 2.27$ and $HV_{WFG9} = HV_{WFG9} + 0.22$.

It is clear from the figure that the WFG problems can
generally be divided into three categories according to the trend of their HV trajectories. The first category corresponds to the problems where the HV trajectory rises with the parameter. The remaining problems are the last category in which HV reaches the best when $SR$ takes some middle values. This indicates that a rare occurrence of the SBS variation is suitable for WFG3 and WFG4, and a frequent one is more acceptable for WFG2 and WFG5. In fact, given that both algorithms (NSGA-II+SBS and NSGA-II) perform fairly well on WFG3 and WFG4 but have difficulties in obtaining a well-converged and well-distributed solution set on WFG2 and WFG5, a low $SR$ value may be advisable for some relatively simple functions while a high $SR$ value seems more useful for intractable problems. In addition, for an unknown optimization problem, the interval $[0.4, 0.6]$ may be more reliable since the HV of the algorithm with $SR$ set in this interval is acceptable on most test problems, as shown in Fig. 11.

In order to further understand the effect of $SR$, Figure 12 shows the “success rate” of individuals generated by SBS that have been inserted into the population during the evolutionary process on WFG3 when $SR = 0$ (denoted by dashed line). Black square denotes the average number of solutions generated by SBS, and hollow circle denotes the average number of solutions inserted into the population.

Fig. 12. Probability that a solution generated by SBS has been inserted into the population during the evolutionary process across the 30 runs of NSGA-II+SBS on WFG3 when $SR = 0$ (denoted by dashed line). Black square denotes the average number of solutions generated by SBS, and hollow circle denotes the average number of solutions inserted into the population.

WFG5 and WFG6 when $MiP = 0.5$ and 0.6, respectively; and for WFG3, WFG4, WFG7, and WFG9 when $MiP = 0.8, 0.6, 0.7$, and 0.8, respectively. Consequently, setting $MiP$ within the range of $[0.4, 0.8]$ seems reliable for an unknown optimization problem.

**Result summary**—High sensitivity of the algorithm to the parameters $SR$ and $MiP$ demands a proper setting of them in SBS. Our experiments suggest that the range of $[0.4, 0.6]$ and $[0.4, 0.8]$ may be reliable for $SR$ and $MiP$, respectively, for an unknown optimization problem. And a median of them (i.e., $SR = 0.5$ and $MiP = 0.6$) may be considered as a first attempt. Furthermore, for some relatively easy problems, a lower $SR$ is recommended, while for some difficult ones, a higher $SR$ may be more suitable.

### V. COMPARISON WITH THREE VARIATION OPERATORS

The primary purpose of our study is to introduce a variation operator to improve the performance of EMO algorithms. For a better validation, we compare SBS with three representative variation operators—opposition-based learning (OBL) [43], jumping gene (JG) [4], and Mühlenbein’s mutation [38].

The OBL operator proposed by Tizhoosh [53] has been found to be competitive in many fields. The main idea of OBL is the simultaneous consideration of an estimate and its corresponding opposite estimate in order to achieve a better approximation for the candidate solution in the population. Let $x_i$ be the value of an individual in the $i$th dimensional decision space ($x_i \in [a_i, b_i]$, $i = 1, 2, ..., D$), and the opposite value of $x_i$ can be calculated as follows:

$$x'_i = a_i + b_i - x_i$$

Similar to the SBS variation, OBL shrinks the search space and the concerned interval $[a_i, b_i]$ is determined by the boundary of the current evolutionary population. However, one important difference is that the place of newly-generated individuals is fixed in OBL, while in SBS newly-generated individuals jump among different segments and vary over the current search space.

JG, which emulates the gene transposition in the genome, is a variation operator to improve the performance of EAs in multiobjective optimization. Inspired from the “jumping gene” phenomenon in biological evolution, the JG operator introduces two types of transpositions: 1) cut-and-paste transposition and 2) copy-and-paste transposition, to exchange information between individuals. JG can be implemented in both binary-valued and real-valued scenarios [4], [39], and the combination of JG with NSGA-II has been found to achieve a good search performance. Further details of the JG variation can be found in [4], [39].

Mühlenbein’s mutation is a mutation operator for real-valued problems. It is able to generate points in the hypercube with the center at a point, defined by $x_i = x_i + range_i \times \gamma$, where $range_i$ defines the mutation range and is normally set to $0.1 \times (b_i - a_i)$, the $+$ or $-$ sign is chosen with a probability of 0.5, and $\gamma$ is defined as follows:

$$\gamma = \sum_{k=0}^{15} \alpha_k 2^{-k}$$


where \( \alpha_i \in \{0, 1\} \) is randomly generated with \( p(\alpha_i = 1) = 1/16 \). In this mutation operator, the probability parameter \( p(\alpha_i = 1) \) being set to 1/16 indicates that on average there will be just one \( \alpha_i \) with value 1. In addition, from the settings of parameter \( k \) in Eq. (9) (i.e., \( k = 0, \ldots, 15 \)), new generated points are located much more often in the neighborhood of \( x_i \) and the minimum possible proximity can be up to a precision of \( \text{range}_i \times 2^{-15} \). Mühlenbein’s mutation is a discrete variation operator, which can be considered as searching among different parts (segments) in the concerned region [18]. However, it has two key differences from the SBS operator: 1) Mühlenbein’s mutation generates points more often in the neighborhood of a given point, and 2) the concerned region in Mühlenbein’s mutation is only a fraction of the whole search space. In addition, it is necessary to point out that despite being a general genetic operator in classical EAs [32], Mühlenbein’s mutation, due to its similarity to SBS, is used here as an auxiliary variation operator to be integrated into EAs to serve the purpose of comparative study.

In order to ensure a fair competition between SBS and the other three operators, the OBL, JG, and Mühlenbein’s mutation operators adopt the evolutionary frame of SBS, and also only act on the nondominated individuals which have been eliminated by the environmental selection (i.e., Algorithm 1 remains unchanged, while Steps 2–4 and Steps 7–24 of Algorithm 2 are replaced by the other operators). NSGA-II is also considered as the test algorithm, and the new three hybrid algorithms are denoted as NSGA-II+OBL, NSGA-II+JG, and NSGA-II+MM, respectively. The parameters used in the peer operators are set according to the recommendation in their original papers, and the common parameters of the four algorithms are kept unchanged.

Table XIII gives the HV results of the peer operators on all the 36 test problems, where the results of NSGA-II+SBS from Table IV are also repeated in this table for the convenience of comparison. Clearly, NSGA-II+SBS considerably outperforms the other three operators, and achieves the best HV value in 18 out of the 36 problems. NSGA-II+OBL, NSGA-II+MM, and NSGA-II+JG perform the best in 3, 8, and 7 out of the 36 problems, respectively. Additionally, for the majority of problems on which NSGA-II+SBS outperforms its competitors, the results have statistical significance (25 out of 32 problems for NSGA-II+OBL, 13 out of 21 problems for NSGA-II+MM, and 19 out of 27 problems for NSGA-II+JG).

In order to further understand the performance difference among these operators, we show their “success rate” during the evolutionary process. Specifically, we show, as a function of the number of evaluations, the probability that a solution generated by SBS has been inserted into the population across the 30 runs of the four algorithms on WFG1 in Fig. 14. Clearly, the success rate of NSGA-II+SBS is higher than that of the other three algorithms, which means that SBS is more efficient to generate “good” solutions that can contribute their value in the evolutionary process.

Result summary—From the above experiments, SBS has shown a clear advantage over the other three variation operators in multiobjective optimization. Hybridized with SBS, NSGA-II can achieve a better HV value than hybridized with OBL, JG, and Mühlenbein’s mutation on half of the tested problems. This means that SBS is a promising operator in enhancing the search ability of EMO algorithms.

VI. Conclusions

This paper presented a variation operator, segment-based search (SBS), to improve the performance of EMO algorithms on continuous MOPs. SBS divides the search space into many small segments according to the current evolutionary status, and performs the macro-jumping and micro-jumping operations inside them to guide the information exchange among “good” individuals. Moreover, the proposed SBS operator is activated when general genetic operators of an algorithm do not work well, thus providing a new power for the algorithm to search towards the desired direction.

Systematic experiments have been carried out by providing an extensive comparison on a set of 36 test problems. From the experimental results, it can be observed that an EMO algorithm with the integration of SBS can improve its performance for MOPs. The new algorithm also outperforms its original version when the number of decision variables shifts. A comparative study on the BRS strategy of SBS reveals that the boundary relaxation is very important to enhance the search ability in multiobjective optimization. In addition, the effect of parameters \( SR \) and \( MiP \) was investigated, and their ranges, \([0.4, 0.6]\) for \( SR \) and \([0.4, 0.8]\) for \( MiP \), are recommended for an unknown optimization problem. Finally, a comparison with three representative variation operators confirmed the competitiveness of the SBS operator in dealing with MOPs.

Segment-based search is a new concept as a variation operator. Further studies are still required to investigate its benefits and limitations. Possible directions of future work include the adaptive setting of the SBS rate and micro-jumping proportion, the investigation of SBS for other algorithms (e.g.,
TABLE XIII
HV results (mean and SD) of NSGA-II+SBS, NSGA-II+OBL, NSGA-II+MM, and NSGA-II+JG, where the best mean among the algorithms for each case is highlighted in boldface.

<table>
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<tr>
<th>Problem</th>
<th>NSGA-II+SBS</th>
<th>NSGA-II+OBL</th>
<th>NSGA-II+MM</th>
<th>NSGA-II+JG</th>
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“*” indicates that the p-value of 58 degrees of freedom is significant at a 0.05 level of significance by a two-tailed t-test for NSGA-II+SBS against its competitor.

Fig. 14. Probability of a solution generated by SBS has been inserted into the population during the evolutionary process across the 30 runs of the considered variation operators (SBS, OBL, JG, and Mühlenbein’s mutation) on WFG1 (denoted by dashed line). Black square denotes the average number of solutions generated by SBS, and hollow circle denotes the average number of solutions inserted into the population.

the Breeder GA), and its improvement for solving MOPs with the Pareto set having a complex shape in the decision space, such as the UF and CF test problems.

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REFERENCES

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