# Population Evolvability: Dynamic Fitness Landscape Analysis for Population-Based Metaheuristic Algorithms

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Abstract—Fitness landscape analysis (FLA) is an important approach for studying how hard problems are for metaheuristic algorithms to solve. Static FLA focuses on extracting the properties of a problem and does not consider any information about the optimization algorithms; thus, it is not adequate for indicating whether a particular algorithm is suitable for solving a problem. By contrast, dynamic FLA considers the behavior of algorithms in combination with the properties of an optimization problem to determine the effectiveness of a given algorithm for solving that problem. However, previous dynamic FLA approaches are all individually based and lack statistical significance. In this paper, the concept of population evolvability is presented, as an extension of dynamic FLA, to quantify the effectiveness of population-based metaheuristic algorithms for solving a given problem. Specifically, two measures of population evolvability are defined that describe the probability that a population will obtain improved solutions to a problem and its ability to do so. Then, a combined measure is derived from these two measures to represent the overall population evolvability. Subsequently, the significance and validity of the proposed measures are investigated through analytical and experimental studies. Finally, the utility of the proposed measures is illustrated in an application of algorithm selection for black-box optimization problems. High accuracy in selecting the best algorithm is observed in a statistical analysis, with a low computational cost in terms of fitness evaluations.

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Index Terms—Algorithm selection task, dynamic fitness landscape analysis, population evolvability, population-based metaheuristic algorithms.

#### I. INTRODUCTION

THE CONCEPT of fitness landscapes originated from theoretical biology more than 80 years ago [1]. In the evolutionary computation community, the notion of fitness landscapes is an attempt to draw an analogy with real landscapes to gain a better understanding of how and where algorithms operate [2]. Fitness landscape analysis (FLA) refers to a collection of data-driven techniques for extracting descriptive or numerical measures related to the properties of fitness landscapes. Hence, the purpose of FLA is to obtain knowledge about a given optimization problem. The most common application of FLA is to gain insights into the characteristics of optimization problems that are critical for algorithm performance [3].

Several descriptive characteristics have been proposed in earlier studies of FLA. For example, the modality, i.e., the distribution of local optima and plateaus, is an intuitive indicator of the hardness of a problem, although the intuitive notion that a multimodal landscape is more difficult than a unimodal landscape is not always true [4], [5]. Basins of attraction are defined with respect to local optima, whose shapes and distributions have been found to be correlated with the computational cost [6]. Fitness barriers are derived from the two concepts discussed above and are represented as a tree-like structure. Two parameters, i.e., the depth and difficulty determined from the barrier tree, play an important role in analysing the convergence speed of simulated annealing [7].

Smoothness, ruggedness and neutrality are the most frequently used measures in studies of FLA. The information provided by these three measures is useful, as in a smooth landscape, an optimization algorithm can easily generate new, fitter observations from its offspring, whereas this is not the case in neutral landscapes, where the directional information contained in the fitness information is zero, or in rugged landscapes, where the fitness values fluctuate rapidly [8]. Several numerical measures have been proposed to describe the degree of ruggedness, such as the autocorrelation function and the correlation length [9], [10]. Neutral walks, which can proceed only within a neutral area, are used to explore neutral areas and record the accumulated distance from the starting point; thus, the maximum accumulated distance can be utilized as an indicator of neutrality [11]. Other measures, such as the

number of neighboring candidate solutions with nearly equal fitness and the number of distinct fitness values surrounding a neutral area, have also been proposed to represent the degree of neutrality [12].

Other methods related to FLA have also been widely adopted. The fitness distance correlation, i.e., the correlation between fitness values and their distances from the global optimum, is designed to provide a global view of the fitness landscape [13]. In addition, exploratory landscape analysis (ELA) approaches, proposed by Mersmann *et al.* [14], [15] and extended by Muñoz *et al.* [16], attempt to extract highlevel and low-level features of fitness landscapes to enable the classification of optimization problems into several categories.

The ultimate goal of FLA is to discover informative relations between properties of fitness landscapes and algorithm performance. Some studies have made considerable progress in this direction. For instance, empirical hardness models were built by Leyton-Brown *et al.* [17], in the context of combinatorial optimization, to predict the run time of an algorithm for a given problem instance. For numerical optimization, a machine learning model was employed by Bischl *et al.* [18] based on the low-level features extracted via the ELA method to predict well-performing algorithms within a given portfolio. Similarly, a meta-learning regression model was used by Muñoz *et al.* [19] based on five existing landscape features to select the proper parameter configurations for CMA-ES [20] on their test problem instances.

The aforementioned techniques all belong to the realm of static FLA, in which measures are extracted from optimization problems without considering any information about the optimization algorithms to be applied. However, the relation between these static measures and algorithm performance is not reliable [21]. Some problems that are judged as plausibly complicated based on their static FLA measures may be easy to solve for certain algorithms, whereas some seemingly easy problems may turn out to actually be hard. Therefore, some researchers have borrowed the notion of evolvability from evolutionary biology to define measures that consider the behavior of algorithms in combination with the properties of optimization problems, in an approach known as dynamic FLA.

In the evolutionary computation community, Altenberg [22] defined evolvability as "the ability of a population to produce variants fitter than any yet existing." It has been demonstrated by Turney [23] that increasing evolvability is regarded as a desirable large-scale trend for evolutionary heuristics; thus, the measures of dynamic FLA are expected to be correlated with algorithm performance. Fitness evolvability portraits were defined by Smith et al. [24] to describe how these measures are related to both ruggedness and neutrality. The negative slope coefficient [25] and the accumulated escape probability (aep) [26] are two measures that indicate the hardness of a problem by measuring the correlation between consecutive solutions produced by the mutation operation of an algorithm. Notably, the existing measures of evolvability are all based on randomly sampled single candidate solutions in the feasible solution space; thus, they are not suitable for analysing the collective behavior of population-based metaheuristic algorithms on a given problem. Additionally, randomness is an

inherent part of the random sampling process, whereas each of these measures provides only a single value as an indicator and thus lacks statistical significance for quantifying this randomness. Therefore, the current dynamic FLA approaches are not applicable in many real optimization scenarios.

Given the above concerns, the question arises of whether it is possible to define measures that are directly correlated with the performance of population-based metaheuristic algorithms on one problem and also have statistical significance. In this paper, we attempt to address this question by presenting the concept of population evolvability, as an extension of dynamic FLA, to measure the effectiveness of population-based metaheuristic algorithms for solving a given optimization problem. Compared with previous studies, the main contributions of this paper are as follows.

- Two measures of population evolvability are formulated, for the first time, to describe the probability that an algorithm will obtain improved solutions to a problem and its ability to do so. Then, a combined measure is derived to represent the overall population evolvability.
- The significance and validity of the proposed measures of population evolvability are investigated through analytical and experimental studies.
- 3) The algorithm selection task for numerical black-box optimization problems is considered as a typical application of the proposed measures, and high accuracy in selecting the best algorithm is observed in a statistical analysis, with a low computational cost in terms of fitness evaluations (FEs).

The remainder of this paper is organized as follows. Section II describes the related work and motivation. Section III is devoted to the explanation of the proposed measures derived from the concept of population evolvability. Section IV presents experimental studies conducted to illustrate the relation between the generational measures and the convergence curve. Section V discusses the algorithm selection task for numerical black-box optimization problems as a typical application of the proposed measures. Further discussions regarding the robustness and computational cost of the proposed approach for the algorithm selection task are presented in Section VI. Finally, Section VII offers the conclusion.

#### II. RELATED WORK AND MOTIVATION

Over the past few decades, FLA has been demonstrated to be a useful tool for analysing the hardness of an optimization problem by extracting its features. Among the available FLA methods, dynamic FLA approaches are most closely related to the measures proposed in this paper, as they are all based on the extraction of features from algorithm-problem pairs. In this section, a brief review of the state of the art in dynamic FLA is first presented. Then, the limitations of current methods are summarized. Finally, the motivation for this paper is introduced.

# A. Fitness Evolvability Portraits

Fitness evolvability portraits were presented by Smith et al. [24] as a means of quantifying the evolvability of

a candidate solution  $\mathbf{x}$ . The authors also described how these portraits relate to both ruggedness and neutrality. Specifically, the fundamental component of a fitness evolvability portrait,  $E_a$ , is defined as follows:

$$E_a = \frac{|N^*(\mathbf{x})|}{|N(\mathbf{x})|} \tag{1}$$

where  $N(\mathbf{x})$  denotes a set of neighbors generated by  $\mathbf{x}$  and  $N^*(\mathbf{x}) = \{\mathbf{c} \mid \mathbf{c} \in N(\mathbf{x}), f(\mathbf{c}) \geq f(\mathbf{x})\}$  denotes a set of nondeleterious neighbors, in which  $f(\cdot)$  denotes a fitness function.  $E_a$  represents the number of nondeleterious neighbors divided by the total number of neighbors; thus, it indicates the probability of nondeleterious mutations.

#### B. Fitness Clouds and the Negative Slope Coefficient

A fitness cloud, as proposed by Philippe *et al.* [27], is a scatter plot that visualizes parent fitness versus offspring fitness. In this approach, the authors assumed that every solution in the search space had only one unique neighbor, which was determined using a search heuristic. The fitness of the neighbor of a solution  $\mathbf{x}$  was called the "bordering fitness,"  $b(\mathbf{x})$ . Then, a fitness cloud could be obtained by constructing a scatter plot from pairs of fitness  $(f(\mathbf{x}))$  and bordering fitness  $(b(\mathbf{x}))$  values. The sets of the minimum, mean, and maximum bordering fitness for every solution were defined as  $FC_{\min}$ ,  $FC_{\max}$ , and  $FC_{\max}$ , respectively, and the intersections of these sets with the diagonal line (i.e., b = f) formed the boundaries between regions of solutions that were strictly advantageous, averagely advantageous, averagely deleterious, and strictly deleterious.

Based on the fitness cloud concept, the authors extracted a singular measure, called the negative slope coefficient (nsc), as an indicator of problem hardness [25]. To calculate nsc, a set of fitness (f) and bordering fitness (b) pairs was divided into consecutive groups according to their fitness values. Then, the mean fitness and the mean bordering fitness within each group, i.e.,  $(\overline{f_i}, \overline{b_i})$ , were calculated. Finally, nsc was calculated as the sum of all negative slopes, as follows:

$$\operatorname{nsc} = \sum_{i} \min(K_i, 0) \tag{2}$$

where  $K_i = (\overline{b_{i+1}} - \overline{b_i})/(\overline{f_{i+1}} - \overline{f_i})$  is the slope between two adjacent groups. This measure, nsc, can be used as an indicator of problem hardness: a problem is easy if nsc = 0, whereas the problem is difficult if nsc < 0, and the magnitude of nsc quantifies the difficulty. As mentioned by the authors, nsc lacks a normalization term, so the results are not comparable across different problems.

# C. Fitness-Probability Clouds and the Accumulated Escape Probability

The concept of a fitness-probability cloud (fpc) was proposed by Lu *et al.* [26] for studying evolvability based on the correlation between the fitnesses of individuals and their escape probabilities. Regarding the escape probability (or escape rate), it was first introduced by Merz [28] to quantify

the number of steps required to escape from a local optimum. Thus, the escape probability,  $p(f_i)$ , is formulated as follows:

$$p(f_i) = \frac{1}{S_i} \tag{3}$$

where  $S_i$  represents the average number of steps required to find an improving move starting from a solution whose fitness value is  $f_i$ . The definition of the escape probability can be extended to a set of fitness values to formulate the average escape probability,  $p_i$ 

$$p_i = \frac{\sum_{f_j \in C_i} p(f_j)}{|C_i|} \tag{4}$$

where  $C_i = \{f_j \mid f_j \geq f_i\}$ . Then the fpc is defined as fpc =  $\{(f_0, p_0), (f_1, p_1), \dots, (f_L, p_L)\}$ .

A numerical measure called the aep was derived from the fpc concept and formulated as follows:

$$aep = \frac{\sum_{f_i \in F} p_i}{|F|} \tag{5}$$

where  $F = \{f_0, f_1, \dots, f_L \mid f_0 < f_1 < \dots < f_L\}$ . aep indicates the hardness of a problem as follows: the larger the value of aep is, the higher the evolvability is, and the easier the problem should be.

#### D. Limitations of Previous Works and Our Motivation

The previous proposed measures of evolvability are all based on randomly sampled single candidate solutions; thus, they are not suitable for directly analysing the collective behavior of population-based metaheuristic algorithms on a given problem. Moreover, each of them provides only a single value as an indicator and thus lacks statistical significance for quantifying the randomness which is an inherent part of the random sampling process. Therefore, in this paper, we wish to develop measures that quantify the evolvability related to sampled populations while also considering the stochastic nature of the sampling process.

The motivation for this paper is as follows: in nature, evolution is correlated with the collective behavior of species, which is controlled by natural selection [29]. Furthermore, the evolutionary rate of a species may change under different circumstances. For instance, as described by Ayala [30], in small or isolated environments such as small islands or remote valleys, evolution can be effectively accelerated. Therefore, we argue that the population evolvability is closely related to two aspects, i.e., the probability of evolution and the ability to evolve. The former represents whether the population can evolve to a fitter state. The latter expresses the degree of that evolution if it occurs.

#### III. MEASURES OF POPULATION EVOLVABILITY

# A. Preliminaries

1) Success Probability: For the convergence analysis of evolution strategies (ESs), the success probability (denoted by  $p_s$  hereafter) was introduced by Rechenberg [31] as part of the famous 1/5th rule for investigating the performance of the

(1+1) ES [32].  $p_s$  is defined as the probability of an offspring replacing its parent. Since there are only one parent and one offspring in the (1+1) ES, the offspring will replace its parent if its fitness value is better than its parent's. Notably, the idea of  $p_s$  is similar to that of the metric  $E_a$  and that of the escape probability  $p(f_i)$  defined in (1) and (3), respectively, as they are all based on an estimation of the probability of successful mutations for single individuals. Specifically,  $p_s$  is measured over a sliding window consisting of a fixed number of the most recent iterations, whereas  $E_a$  and  $p(f_i)$  are estimated based on a fixed number of neighbors generated by an individual.

2) Convergence Velocity and Convergence Reliability: Convergence velocity and convergence reliability analyses have achieved considerable progress in theoretical research on evolutionary algorithms. According to [33], the convergence velocity is a local measure of improvement from one iteration to the next, where the improvement can be measured in terms of either the expected quality gain,  $\varphi$ , or the expected change in the distance to the global optimum, d. Concretely,  $\varphi$  and d are formulated as follows:

$$\varphi = E[f(\mathbf{x}_{t+1}) - f(\mathbf{x}_t)]$$

$$d = E[\|\mathbf{x}^* - \mathbf{x}_t\| - \|\mathbf{x}^* - \mathbf{x}_{t+1}\|]$$
(6)

where  $\mathbf{x}^*$  denotes the global optimum and  $\mathbf{x}_t$  represents the best or a representative individual of the population at generation t. A convergence reliability analysis focuses on the global convergence of an algorithm within an unlimited run time, that is

$$\lim_{t \to \infty} p(\mathbf{x}^* \in P(t)) = 1 \tag{7}$$

where P(t) denotes the population at generation t and  $p(\cdot)$  is the probability of an event.

Regarding the above two preliminaries, the success probability estimates the probability of successful mutations, whereas the convergence velocity and convergence reliability measure the convergence ability of evolutionary algorithms. These concepts are closely related to the two aspects of evolution that we wish to assess, i.e., the probability of and ability for evolution, respectively, and thus provide inspiration for formulating our measures of population evolvability.

#### B. Evolutionary Probability of Population

The concept of the probability of successful mutations of an individual is expected to be extended to that of the probability of the successful evolution of a population. As we know, metaheuristic algorithms typically include various operators and strategies, such as recombination, mutation, selection, and self-adaption; consequently, the neighbors (one-step child populations) are the products of comprehensive effects of multiple operations on their common parent, not merely the results of a mutation operation, as considered in previous studies. Therefore, we need a new measure to reflect the evolutionary probability of a population (epp).

Definition 1: Given a sampled population  $P_i$  and a set of its generated neighbors (one-step child populations)

 $N(P_i) = \{P_{i1}, \dots, P_{iM} \mid M \ge 1\}$ , define the epp to be

$$epp(P_i) = \frac{|N^+(P_i)|}{|N(P_i)|}$$
 (8)

where  $N^+(P_i)$  represents the set of evolved neighbors, i.e.,  $N^+(P_i) = \{P_{ij} \mid P_{ij} \in N(P_i), f^b(P_{ij}) < f^b(P_i)\}$  for a minimization problem, in which  $j = 1, \ldots, |N^+(P_i)|$  represents the index of evolved neighbors, and  $f^b(P)$  denotes the best fitness value found in a population P.

Although we focus only on the best individual in the population and its generated neighbors, Definition 1 truly concerns the collective behavior of the entire population. Moreover, through this definition, we can investigate the population evolvability for a large variety of population-based metaheuristic algorithms based on a unified concept, i.e., the probability that a population will evolve from its *current state*. Notably, the meaning of the current state may differ for different algorithms. To be more concrete, for algorithms that have no self-adaptive parameters, such as the real-coded GA [34], the current state of a population comprises the genotype and phenotype of the population and the global and fixed parameters of the algorithm, whereas for algorithms that do have selfadaptive parameters, such as CMA-ES [20], the current values of these adaptive parameters should also be included in the current state. Note that the range of epp is [0, 1], as it is an estimation of a probability.

#### C. Evolutionary Ability of Population

Evolutionary abilities of populations in different states are diverse. For metaheuristic algorithms, it is far from sufficient to use the convergence velocity  $\varphi$  defined in (6) as a measure of evolutionary ability. The reasons are as follows: first, the convergence velocity is a local metric and consequently cannot reflect the global behavior of the algorithm, yet this kind of global behavior is vital to metaheuristic algorithms for exploring the solution space, and second, the convergence velocity mainly focuses on the absolute changes between fitness values, which will become increasingly smaller while approaching convergence, making it unable to represent the evolution ability near convergence. Therefore, we need a new measure to reflect the evolutionary ability of an algorithm throughout the entire optimization process.

Definition 2: Given  $P_i$  and  $N^+(P_i)$  that are the same as in Definition 1, define the evolutionary ability of a population (eap) to be

$$\operatorname{eap}(P_i) = \begin{cases} \frac{\sum_{P_{ij} \in N^+(P_i)} \frac{|f^b(P_i) - f^b(P_{ij})| / NP}{\sigma(f(P_i))}}{|N^+(P_i)|}, & |N^+(P_i)| \ge 1\\ 0, & |N^+(P_i)| = 0 \end{cases}$$
(9)

where NP represents the size of the population  $P_i$ , and  $\sigma(f(P_i))$  represents the standard deviation of the fitness values of the population  $P_i$ .

The meaning of Definition 2 can be interpreted as the average eap calculated over its evolved neighbors, where the

evolutionary ability is expressed as follows:

$$\frac{|f^b(P_i) - f^b(P_{ij})|/NP}{\sigma(f(P_i))}.$$
 (10)

In (10), the numerator is the absolute improvement in the best fitness value between the population  $P_i$  and its evolved neighbor  $P_{ij}$ . The reason for dividing by NP is to obtain equitable improvements for populations of different sizes. As for the denominator, it is the standard deviation of the fitness values of  $P_i$ , which is a distance-based diversity measure with respect to the population phenotype, as proposed in [35]. For this equation, on the one hand, if the denominator remains constant, the larger the numerator is, the greater the evolutionary ability will be; that is to say, if the diversity measures of different populations are equal, then the population that can achieve the highest absolute fitness improvement is the one that has the greatest evolutionary ability. On the other hand, if the numerator remains constant, the smaller the denominator is, the greater the evolutionary ability will be; that is to say, if the absolute fitness improvements of different populations are the same, then the population with the lowest diversity is considered to have the greatest evolutionary ability. In both situations, this equation is able to provide a reasonable explanation for the evolutionary ability. Notably, if  $\sigma(f(P_i))$ is equal or infinitely approaching to zero and the numerator is nonzero, then eap could be infinitely large. To avoid this situation, we set  $\sigma(f(P_i)) = \varepsilon$  when  $\sigma(f(P_i)) < \varepsilon$  (where  $\varepsilon$ is a small constant). In this paper,  $\varepsilon$  is set to 1e-8.

Considering the optimization process of metaheuristic algorithms, large fitness improvements and high diversity are the features of earlier stages, whereas relatively smaller fitness improvements and lower diversity are the ordinary states of later stages. We expect that eap can serve as a measure of evolutionary ability throughout the entire process of optimization, although we do not exclude the possibility that measures of other forms might achieve the same purpose. Finally, it should be noted that the range of eap is  $[0, +\infty)$ .

# D. Evolvability of Population

From an algorithmic perspective, a problem is hard either if it is hard for evolution to occur (i.e., the evolutionary probability is low) or if the degree of evolution is small (i.e., the evolutionary ability is low). Thus, we wish to consider both phenomena simultaneously to formulate a combined measure of problem hardness for population-based metaheuristic algorithms. Certainly, there are various possible ways to combine these two aspects; in particular, we adopt a concise and meaningful one, namely, multiplying them together, to reflect the overall population evolvability.

Definition 3: Given epp and eap, define the evolvability of a population (evp) to be

$$\exp(P_i) = \exp(P_i) \times \exp(P_i)$$

$$= \begin{cases} \frac{\sum_{P_{ij} \in N^+(P_i)} \frac{|f^b(P_i) - f^b(P_{ij})|/NP}{\sigma(f(P_i))}}{|N(P_i)|}, & |N^+(P_i)| \ge 1\\ 0, & |N^+(P_i)| = 0. \end{cases}$$
(11)

Definition 3 represents the average evolutionary ability over the entire set of generated neighbors, where the evolution abilities of unevolved neighbors are considered to be 0. In addition, because epp is defined as the probability of evolution and eap is a measure of evolutionary ability, evp actually represents the expectation of the evolutionary ability for a population  $P_i$ . In particular, when  $P_i$  exists in a given state,  $\exp(P_i)$  reflects the expected evolutionary ability from that state, which is correlated with the hardness of the problem for the algorithm in that state. Clearly, the range of evp is  $[0, +\infty)$ .

#### IV. EXPERIMENTAL STUDIES

This section presents experimental studies conducted to demonstrate the validity of the proposed measures of population evolvability. First, thirteen benchmark functions and a test problem adapted from a real application are introduced. Then, five representative population-based algorithms are described. Finally, the results are presented and analyzed in detail.

#### A. Problem Set

Thirteen traditional numerical problems among the benchmark functions of CEC 2013 [36] are considered first. All of these functions are minimization problems, whose search space lies in  $[-100, 100]^D$  and whose global optimum is randomly generated within  $[-80, 80]^D$ . Note that the fitness value of the global optimum is separately shifted to positive, zero, and negative values to evaluate the proposed measures under various circumstances. In addition, we treat all selected test functions as black-box optimization problems, each with 30 dimensions, i.e., D = 30. Table I lists the details of the selected test functions, where the first five functions are unimodal and the last eight functions are multimodal.

A real-world optimization problem for the design of spreadspectrum radar polyphase codes [37] (denoted by  $f_{\text{radar}}$  hereafter) is also used as a test problem. It can be modeled as a min-max nonlinear and nonconvex optimization problem with continuous variables and numerous local optima, which has been proven to be NP-hard [38]. This problem can be formulated as follows:

global min 
$$f(\mathbf{x}) = \max\{\varphi_1(\mathbf{x}), \dots, \varphi_{2m}(\mathbf{x})\}\$$
  
 $\mathbf{x} = \{(x_1, \dots, x_D) \in \mathbb{R}^D \mid 0 \le x_j \le 2\pi, j = 1, \dots, D\}\$   
 $\varphi_{2i-1}(\mathbf{x}) = \sum_{j=1}^D \cos\left(\sum_{k=|2i-j-1|+1}^j x_k\right), i = 1, \dots, D$   
 $\varphi_{2i}(\mathbf{x}) = 0.5 + \sum_{j=i+1}^D \cos\left(\sum_{k=|2i-j|+1}^j x_k\right), i = 1, \dots, D-1$   
 $\varphi_{m+i}(\mathbf{x}) = -\varphi_i(\mathbf{x}), i = 1, \dots, m; m = 2D-1.$  (12)

The objective of this optimal design problem is to minimize the largest module among the samples of the auto-correlation function, which is correlated with the complex envelope of the compressed radar pulse at the output of an optimal receiver. The surface plot for this problem is shown in Fig. 1, which

Function	Definition	Transformation	$f_i^*$
Sphere Function	$f_1(\mathbf{x}) = \sum_{i=1}^D z_i^2 + f_1^*$	z = x - o	-600
High Conditioned Elliptic Function	$f_2(\mathbf{x}) = \sum_{i=1}^{D} (10^6)^{\frac{i-1}{D-1}} z_i^2 + f_2^*$	$\mathbf{z} = T_{osz}(\mathbf{x} - \mathbf{o})$	-500
Bent Cigar Function	$f_3(\mathbf{x}) = z_1^2 + 10^6 \sum_{i=2}^D z_i^2 + f_3^*$	$\mathbf{z} = T_{asy}^{0.5}(\mathbf{x} - \mathbf{o})$	-400
Discus Function	$f_4(\mathbf{x}) = 10^6 z_1^2 + \sum_{i=2}^D z_i^2 + f_4^*$	$\mathbf{z} = T_{osz}(\mathbf{x} - \mathbf{o})$	-300
Different Powers Function	$f_5(\mathbf{x}) = \sqrt{\sum_{i=1}^{D}  z_i ^{2+4\frac{i-1}{D-1}}} + f_5^*$	z = x - o	-200
Rosenbrock's Function	$f_6(\mathbf{x}) = \sum_{i=1}^{D-1} (100(z_i^2 - z_{i+1})^2 + (z_i - 1)^2) + f_6^*$	$\mathbf{z} = \frac{2.048(\mathbf{x} - \mathbf{o})}{100} + 1$	-100
Schaffers F7 Function	$f_7(\mathbf{x}) = \left(\frac{1}{D-1} \sum_{i=1}^{D-1} (\sqrt{z_i} + \sqrt{z_i} \sin^2(50z_i^{0.2}))\right)^2 + f_7^*$	$z_i = \sqrt{y_i^2 + y_{i+1}^2}$ $\mathbf{y} = \Lambda^{10} T_{asy}^{0.5} (\mathbf{x} - \mathbf{o})$	0
Ackley's Function	$f_8(\mathbf{x}) = -20 \exp\left(-0.2\sqrt{\frac{1}{D}\sum_{i=1}^D z_i^2}\right) - \exp\left(\frac{1}{D}\sum_{i=1}^D \cos(2\pi z_i)\right) + 20 + e + f_8^*$	$\mathbf{z} = \Lambda^{10} T_{asy}^{0.5} (\mathbf{x} - \mathbf{o})$	100
Weierstrass Function	$f_9(\mathbf{x}) = \sum_{i=1}^{D} \left( \sum_{k=0}^{k_{max}} [a^k \cos(2\pi b^k (z_i + 0.5))] \right) - D \sum_{k=0}^{k_{max}} [a^k \cos(2\pi b^k \cdot 0.5))] + f_9^*$	$a = 0.5, b = 3, k_{max} = 20,$ $\mathbf{z} = \Lambda^{10} T_{asy}^{0.5} \left( \frac{0.5(\mathbf{x} - \mathbf{o})}{100} \right)$	200
Griewank's Function	$f_{10}(\mathbf{x}) = \sum_{i=1}^{D} \frac{z_i^2}{4000} - \prod_{i=1}^{D} \cos(\frac{z_i}{\sqrt{i}}) + 1 + f_{10}^*$	$\mathbf{z} = \Lambda^{100} \left( \frac{600(\mathbf{x} - \mathbf{o})}{100} \right)$	300
Rastrigin's Function	$f_{11}(\mathbf{x}) = \sum_{i=1}^{D} (z_i^2 - 10\cos(2\pi z_i) + 10) + f_{11}^*$	$\mathbf{z} = \Lambda^{10} T_{asy}^{0.2} \left( T_{osz} \left( \frac{5.12(\mathbf{x} - \mathbf{o})}{100} \right) \right)$	400
Schwefel's Function	$f_{12}(\mathbf{x}) = 418.9829 \times D - \sum_{i=1}^{D} g(z_i) + f_{12}^*$	$\mathbf{z} = \Lambda^{10} \left( \frac{1000(\mathbf{x} - \mathbf{o})}{100} \right) + 420.9687462275036$	500
Katsuura Function	$f_{13}(\mathbf{x}) = \frac{10}{D^2} \prod_{i=1}^{D} \left( 1 + i \sum_{j=1}^{32} \frac{ 2^j z_i - round(2^j z_j) }{2^j} \right)^{\frac{10}{D^{1,2}}} -$	$\mathbf{z} = \Lambda^{100} \left( \frac{5(\mathbf{x} - \mathbf{o})}{100} \right)$	600

TABLE I
THIRTEEN REVISED TEST FUNCTIONS FROM CEC 2013



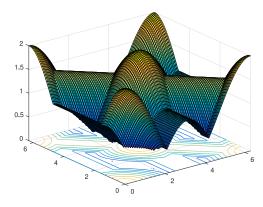


Fig. 1. Surface plot of the spread spectrum radar polyphase code design problem.

illustrates its complexity in D=2. In this paper, the dimensionality of this problem is set to 20 (i.e., D=20), which makes it even more complicated to solve.

# B. Algorithm Settings

Five representative and state-of-the-art population-based metaheuristic algorithms [39], including evolutionary and swarm-based optimization techniques for numerical function optimization, i.e., the real-coded GA [34], CMA-ES [20], CoDE [40], SPSO2011 [41], and ABC [42], are studied. Among them, the real-coded GA was implemented by modifying the "ga" function of the MATLAB Global Optimization Toolbox [43], where the strategy for parent selection is "rank fitness scaling" followed by "stochastic universal sampling"; the crossover and mutation operators are "whole arithmetic" and "uniform," respectively; and survival selection is performed using "elitism." The other four algorithms were all implemented from the source codes provided online by their authors. In addition, the bounded constraint strategy "absorb" [i.e.,  $min(max(\mathbf{x}, lb), ub)$ ] [44] was added to all algorithms except for CMA-ES and SPSO2011, as these algorithms possess specific strategies provided by their authors for handling the search space boundaries. We set the parameters of the real-coded GA, i.e., the crossover rate  $(p_c)$ , mutation rate  $(p_m)$ and elite ratio  $(r_e)$ , to  $p_c = 0.8$ ,  $p_m = 0.05$ , and  $r_e = 0.05$ , respectively. The population sizes for all algorithms were set following the recommendations of their authors; i.e., 100 for the real-coded GA,  $4 + \lfloor 3 + \ln D \rfloor$  for CMA-ES, 30 for CoDE, 40 for SPSO2011, and 50 for ABC.<sup>1</sup>

<sup>1</sup>The source code required to run all of the experiments described in this paper is available at https://github.com/mangwang/TEVC17.

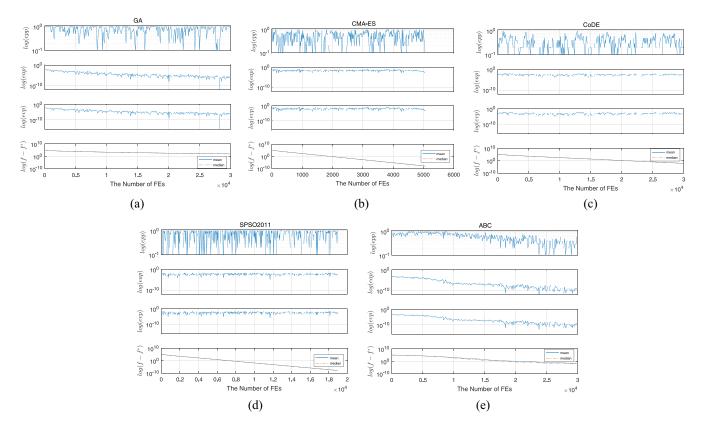


Fig. 2. Relationship between the generational measures and the convergence curve for the sphere function. In each group of four, the top graph shows  $\log(\text{epp})$ , the second graph shows  $\log(\text{eap})$ , the third graph shows  $\log(\text{evp})$ , and the bottom graph shows  $\log(f - f^*)$ . As the y axes of these graphs are on a logarithmic scale, gaps appear in continuous regions where the measures take a value of zero. (a) Sphere/GA. (b) Sphere/CMA-ES. (c) Sphere/CoDE. (d) Sphere/SPSO2011. (e) Sphere/ABC.

# C. Results of Visualization

The relationship between the proposed measures of population evolvability and the convergence curve is visualized for all test problems. In this process, we unified the budget of FEs to  $10^3 \times D$  and assumed the fitness value of the global optimum to be known a priori for better visualization of the convergence curve. Each of the studied algorithms was regularly run on every test problem to generate a sequence of  $P_i$ ; then, the proposed measures of population evolvability were estimated for ten neighbors separately following (8), (9), and (11) with  $f^b(P)$  replaced with  $f_t^b(P)$ , which represents the best fitness value found up through generation t. Each convergence curve plot showed the generational mean and median of 50 convergence curves that were generated by independently running the studied algorithm on the test problem. We initialized the random number generator with a time-dependent seed to ensure the randomness of each run and independence among all runs. Additionally, in accordance with the IEEE Standard for Floating-Point Arithmetic (IEEE 754), the total precision of the double-precision floating-point format is 53 bits, which corresponds to 15-17 significant decimal digits of precision. Therefore, to strike a balance between precision and practicability, we set the threshold to 1e-8and revised the fitness value based on the following strategy: if  $f - \lfloor f \rfloor < 1e - 8$ , then  $f = \lfloor f \rfloor$ . Because of the page limit, we show the results for only three typical test problems, i.e., the sphere function, Schwefel's function, and the spread-spectrum radar polyphase code design problem (called the radar problem hereafter), as these problems represent simple unimodal, complicated multimodal, and NP-hard numerical optimization problems, respectively. The results of the rest of the test problems are included in the supplementary material.

Figs. 2–4 present the visualization results for the application of the studied algorithms to the sphere function, Schwefel's function, and the radar problem. From these figures, several interesting findings can be extracted. First, for an easy optimization problem such as the sphere function, the values of epp for all studied algorithms remain relatively high throughout the entire process until the global optimum is reached (with the floating-point precision correction), whereas the magnitudes of eap for the different algorithms are diverse. Further inspection of the convergence curves reveals that larger magnitudes of eap correspond to faster convergence toward the global optimum. In this respect, CMA-ES has the largest eap magnitudes; thus, it is the fastest algorithm that finds the global optimum. Second, for hard problems such as Schwefel's function and the radar problem, either epp is relatively low [e.g., for CMA-ES on Schwefel's function and for CoDE on the radar problem, as shown in Figs. 3(b) and 4(c), respectively], or eap is relatively low [e.g., for SPSO2011 on Schwefel's function and the radar problem, as shown in Figs. 3(d) and 4(d), respectively]. Such observations illustrate that these algorithms have difficulty evolving toward a fitter region because of either

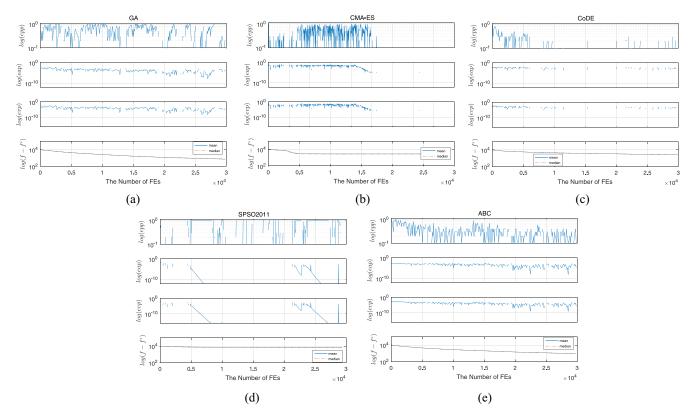


Fig. 3. Relationship between the generational measures and the convergence curve for Schwefel's function. In each group of four, the top graph shows  $\log(\text{epp})$ , the second graph shows  $\log(e_p)$ , the second graph shows  $\log(f-f^*)$ . As the y axes of these graphs are on a logarithmic scale, gaps appear in continuous regions where the measures take a value of zero. (a) Schwefel/GA. (b) Schwefel/CMA-ES. (c) Schwefel/CoDE. (d) Schwefel/SPSO2011. (e) Schwefel/ABC.

a low evolutionary probability or a low evolutionary ability. Moreover, the performances of these algorithms are also not good, as shown in Figs. 3(b) and (d) and 4(c) and (d), which indicates that these two aspects of problem hardness for a population-based metaheuristic algorithm are effectively reflected in the values of epp and eap. As a result, the evp measure is expected to be correlated with the overall problem hardness. In fact, all of the figures illustrate that the values of evp are strongly related to the performances of the studied algorithms, regardless of the test problem. Concretely, if evp remains at a relatively high value, as shown in Fig. 2 for example, the best fitness values obtained by the algorithm in generations will continuously improve. If, however, the evp measure exhibits continuous regions of zero values, as for CMA-ES and SPSO2011 on Schwefel's function [shown in Fig. 3(b) and (d)], then the best fitness values achieved by the algorithm will remain nearly unchanged, and improvement will be difficult to achieve.

# V. ALGORITHM SELECTION TASK FOR BLACK-BOX OPTIMIZATION PROBLEMS—APPLICATION

From the visualization results presented above, the values of evp are expected to be closely related to the problem hardness for metaheuristic algorithms. Therefore, in this section, we utilize the sequence of evp values as an indicator for performing the algorithm selection task for the test problems described in Section IV. First, the algorithm selection task is introduced. Then, the overall framework for algorithm selection using

the evp sequence is explained in detail. Finally, the results achieved using the proposed algorithm selection framework are evaluated through a statistical comparison with the traditional performance evaluation approach, and high accuracy is found to be achieved at a low computational cost.

#### A. Algorithm Selection Task

The algorithm selection task, which was first proposed by Rice [45], consists of four components: 1) the problem space, F; 2) the algorithm space, A; 3) the performance space, Q; and 4) the characteristic space, C. For a given problem instance  $f \in F$ , the algorithm  $\alpha \in A$  that achieves the best performance in Q can be selected based on the extracted characteristics from C. According to this definition, in our case, the values of the evp measure can be viewed as the characteristics obtained via dynamic FLA for population-based metaheuristic algorithms; thus, they are expected to be feasible for the algorithm selection task.

#### B. Overall Framework

Fig. 5 depicts the overall framework for employing the values of  $\exp(P_i)$  for the algorithm selection task, where the input to this framework is a set of candidate algorithms A, and the best algorithm  $\alpha$  is selected as its output. This framework consists of four relevant components.

1) The sampling of the feasible solution space to obtain sampled populations.

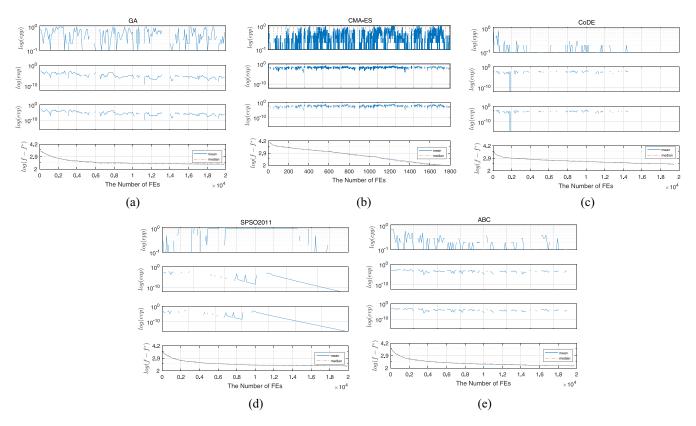


Fig. 4. Relationship between the generational measures and the convergence curve for the radar problem. In each group of four, the top graph shows  $\log(\text{epp})$ , the second graph shows  $\log(\text{eap})$ , the third graph shows  $\log(\text{evp})$ , and the bottom graph shows  $\log(f-f^*)$ . As the y axes of these graphs are on a logarithmic scale, gaps appear in continuous regions where the measures take a value of zero. (a) Radar/GA. (b) Radar/CMA-ES. (c) Radar/CoDE. (d) Radar/SPSO2011. (e) Radar/ABC.

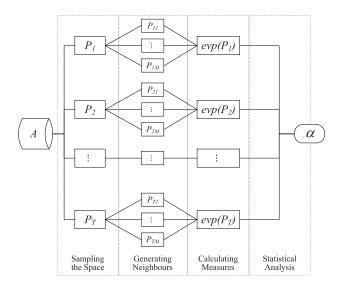


Fig. 5. Overall framework for employing evp for the algorithm selection task.

- 2) The generation of a fixed number of neighbors (one-step child populations).
- 3) The calculation of the  $evp(P_i)$  values.
- A statistical analysis for selecting the appropriate algorithm.

These components are described in detail below.

1) Sampling the Solution Space: In general, sampling should be used to generate populations (i.e.,  $P_1, P_2, \ldots, P_T$ 

shown in Fig. 5) that can represent various states, as the size of the feasible solution space does not permit the consideration of all possible regions. Sampling can be achieved either offline or online. With regard to offline sampling, such as latin hypercube sampling and random sampling [46], the same weights are assigned to all valid search areas, whereas the Metropolis method [47] assigns higher weights to "important" areas. Because not all areas are equally important for metaheuristic algorithms, the Metropolis method is preferred. However, the size and location of the search region sampled by an algorithm may constantly change during optimization. Moreover, some self-adaptive algorithms contain adaptive parameters, and no offline sampling method can obtain information about the search region or the current state of adaptive parameters, which means that such methods are not suitable for our purposes. By contrast, with an online sampling method, the sampling procedure is performed during the execution of the algorithm itself. Specifically, the first population  $P_1$  is uniformly randomly sampled from the entire feasible search space, and subsequent populations are generated throughout the execution of the algorithm until a termination criterion is met. Online sampling methods are able to discover important areas and update adaptive parameters automatically by virtue of the behavior of metaheuristic algorithms, and hence, the online sampling approach is employed in this paper.

When online sampling is adopted, a termination criterion should be established. As different algorithms may have different population sizes, we use FEs as our measure of

computational cost [48], and a budget of FEs is used as the termination criterion. Concretely, the FEs budget for online sampling (denoted by samFEs hereafter) are initially limited to a number much smaller than the maxFEs budget,<sup>2</sup> such as samFEs = 20%maxFEs. Then, for a given optimization problem, a candidate algorithm performs online sampling in the feasible space until the samFEs budget is reached.

2) Generating Neighbors and Calculating Measure Values: Online sampling generates parental populations, and the best fitness value obtained by each algorithm is recorded during the sampling process. As we know, on simple optimization problems such as the sphere function, certain algorithms (e.g., CMA-ES) can rapidly reach the global optimum (with the floating-point precision correction), such that the number of FEs used may be even fewer than samFEs. In this situation, if we were still to calculate the evp values after the convergence of the algorithm to the global optimum, they would be consecutive zeros, which is unreasonable. Therefore, to ensure fair comparisons among the candidate algorithms and in consideration of the fact that the fitness value of the global optimum is usually unknown in real applications, the best fitness value achieved by the best-performing algorithm during online sampling is considered as an approximation of the fitness value of the global optimum. Thus, the number of FEs corresponding to the first hitting time of the algorithm that is the fastest to reach the fitness value of approximate global optimum is considered as the unified budget for subsequent procedures (denoted by uniFEs hereafter), which means that sampled populations that use more FEs than uniFEs are discarded in later procedures. In brief, uniFEs can be less than or equal to samFEs, depending on whether the algorithm can hit the fitness value of the approximate global optimum before reaching the samFEs budget.

For each candidate algorithm, the sampled populations within the *uniFEs* budget are treated as parents to separately generate M neighbors from their current states. All neighbors are randomly and independently generated throughout the algorithm to ensure the generalizability of the proposed framework. Then, each  $\exp(P_i)$  value can be calculated following the procedure shown in Fig. 6, in which, for two consecutive parental populations, i.e.,  $P_i$  and  $P_{i+1}$  (where  $i=1,2,\ldots,T-1$ ),  $P_{i+1}$  is also a neighbor of  $P_i$ ; thus, we need to generate only (M-1) new neighbors to estimate the value of  $\exp(P_i)$ . Fig. 6 also indicates that the minimum value of M is 1, in which case, each  $\exp(P_i)$  value is estimated directly through the online sampling process.

3) Statistical Analysis: For a given optimization problem, each candidate algorithm generates a sequence of  $\exp(P_i)$  values. The sequence sizes for different algorithms may be diverse because of different population sizes. Further investigation of the relations indicates that because of the intrinsic behavior of metaheuristic algorithms, the  $\exp(P_i)$  values in each sequence are dependent and not normally distributed, and all sequences are independent of each other. Moreover, there are usually at least two candidate algorithms, which means that the number

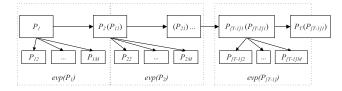


Fig. 6. Procedure for calculating  $\exp(P_i)$ , in which  $P_{ij}$  denotes the neighbors of  $P_i$ . Because of the online sampling procedure,  $P_{i+1}$  is also a neighbor (shown in parentheses) of  $P_i$ .

of sequences is usually greater than two. These properties of the evp sequences satisfy all of the requirements for a non-parametric statistical test, i.e., the Kruskal–Wallis [49] test. In this test, the null hypothesis  $H_0$  is that the distributions of all k sequences are equal, where k is the number of candidate algorithms. Then, the total evp values from all sequences are globally ranked. Let  $N_s$  denotes the total number of evp values across all sequences,  $r_{ij}$  denotes the global rank of evp value j from sequence i,  $n_i$  denotes the number of evp values in sequence i, and  $R_i = \sum_{j=1}^{n_i} r_{ij}$  denotes the sum of the ranks from the ith sequence, then  $N_s = \sum_{i=1}^k n_i$ , and the test statistic is calculated in the following form:

$$H = \frac{12}{N_s(N_s+1)} \sum_{i=1}^k \frac{R_i^2}{n_i} - 3(N_s+1).$$
 (13)

Any tied values are assigned the average of their original ranks, and a correction for ties can be applied by dividing the H value calculated in (13) by  $1 - \sum_{g=1}^{G} (t_g^3 - t_g)/(N_s^3 - N_s)$ , where G represents the number of groupings of different tied ranks and  $t_g$  is the number of tied ranks within group g. The p-value of the Kruskal–Wallis test can be approximated from the  $\chi^2$  distribution with (k-1) degrees of freedom. If the p-value is less than the preset significance level (such as 0.05), a post hoc procedure can be used for multiple comparisons between pairs of sequences to obtain stochastic dominance [50]. In this paper, we adopt the Conover–Iman [51] test as the post hoc procedure because it can directly use the ranks generated from the Kruskal–Wallis test. Its test statistic is formulated as follows:

$$S^{2} = \frac{1}{N_{s} - 1} \left[ \sum_{i=1}^{k} \sum_{j=1}^{n_{i}} r_{ij}^{2} - N_{s} \frac{(N_{s} + 1)^{2}}{4} \right]$$
$$\left| \overline{R_{i}} - \overline{R_{j}} \right| > t_{1-\alpha} \sqrt{S^{2} \frac{N_{s} - 1 - H}{N_{s} - k}} \sqrt{\frac{1}{n_{i}} + \frac{1}{n_{j}}}$$
(14)

where  $\overline{R_i} = R_i/n_i$  represents the average rank in sequence i and t represents Student's t-distribution. Then, the p-value can be calculated from the upper  $\alpha$  quantile of the t-distribution with  $(N_s - k)$  degrees of freedom and adjusted by means of the Holm [52] procedure to revise the family wise error rate (FWER), which represents the probability that one or more false conclusions will be drawn among all hypotheses when performing multiple pair-wise tests. Eventually, the best algorithm or a set of equally best algorithms can be selected according to the results of the statistical tests described above.

<sup>&</sup>lt;sup>2</sup>The maxFEs budget depends on the optimization problem and is usually set based on either experience or typical recommendations from others.

TABLE II								
COMPARISON OF	COMPARISON OF THE RESULTS ACHIEVED USING THE PROPOSED APPROACH AND THE							
PERFORMANC	PERFORMANCE EVALUATION APPROACH FOR THE ALGORITHM SELECTION TASK							
GA	CMA-ES	CoDE	SPSO2011					
201	407 <sup>†</sup>	182	228					

Function	Measure	GA	CMA-ES	CoDE	SPSO2011	ABC
r.	avgEvpRank	201	407 <sup>†</sup>	182	228	88
$f_1$	performEval	-6.0000E+02(3.54E-04)	$-6.0000E+02(0.00E+00)^{\dagger}$	$-6.0000E+02(0.00E+00)^{\dagger}$	$-6.0000E+02(0.00E+00)^{\dagger}$	-6.0000E+02 $(0.00E+00)^{\dagger}$
£_	avgEvpRank	916	3479 <sup>†</sup>	1870	1951	817
$f_2$	performEval	-4.9988E+02(1.12E-01)	$-5.0000E+02(0.00E+00)^{\dagger}$	-5.0000E+02 $(0.00E+00)^{\dagger}$	2.1386E+05(6.98E+04)	-5.0000E+02 $(0.00E+00)^{\dagger}$
r	avgEvpRank	320	$1202^{\dagger}$	698	749	235
$f_3$	performEval	-9.9465E+01(2.86E+02)	$-4.0000E+02(0.00E+00)^{\dagger}$	$-4.0000E+02(0.00E+00)^{\dagger}$	1.8478E+03(2.63E+03)	-4.0000E+02 $(0.00E+00)^{\dagger}$
£	avgEvpRank	1839	2118	$2648^{\dagger}$	1174	1594
$f_4$	performEval	-3.0000E+02(4.52E-04)	$-3.0000E+02(0.00E+00)^{\dagger}$	$-3.0000E+02(0.00E+00)^{\dagger}$	1.7126E+03(6.77E+02)	$-3.0000E+02(0.00E+00)^{\dagger}$
r	avgEvpRank	1332	4797 <sup>†</sup>	2905	3659	1103
$f_5$	performEval	-2.0000E+02(2.04E-03)	$-2.0000E+02(0.00E+00)^{\dagger}$	$-2.0000E+02(0.00E+00)^{\dagger}$	-2.0000E+02(3.98E-05)	$-2.0000E+02(0.00E+00)^{\dagger}$
£	avgEvpRank	898	$3017^{\dagger}$	2086	1455	960
$f_6$	performEval	-9.1746E+01(2.17E+01)	$-9.8563E+01(1.01E+01)^{\dagger}$	$-9.7931E+01(1.09E+01)^{\dagger}$	-3.0970E+01(3.40E+01)	-9.9995E+01(5.33E-03)
$f_7$	avgEvpRank	4181	3173	5397 <sup>†</sup>	4582	3889
	performEval	1.4193E-01(1.05E-01)	1.7880E+01(1.38E+01)	$0.0000E+00(0.00E+00)^{\dagger}$	4.6581E+01(1.52E+01)	$0.0000E+00(0.00E+00)^{\dagger}$
$f_8$	avgEvpRank	5436 <sup>†</sup>	2991	4657	4220	$5334^{\dagger}$
	performEval	$1.2000E+02(2.33E-04)^{\dagger}$	1.2024E+02(1.23E-01)	1.2000E+02(3.00E-03)	1.2075E+02(8.62E-02)	1.2000E+02(8.26E-04)
r	avgEvpRank	5475	3073	4234	4321	5913 <sup>†</sup>
$f_9$	performEval	2.0006E+02(2.30E-02)	2.0445E+02(2.54E+00)	2.0050E+02(4.25E-01)	2.1400E+02(3.52E+00)	$2.0000E+02(2.44E-05)^{\dagger}$
avgEvpRa	avgEvpRank	4565	2710	5583 <sup>†</sup>	5639 <sup>†</sup>	4467
$f_{10}$	performEval	3.0008E+02(4.94E-02)	3.0002E+02(1.16E-02)	$3.0000E+02(0.00E+00)^{\dagger}$	3.0021E+02(1.15E-01)	$3.0000E+02(0.00E+00)^{\dagger}$
£	avgEvpRank	5441	2758	4823	4937	5832 <sup>†</sup>
$f_{11}$	performEval	4.0000E+02(2.76E-03)	4.5015E+02(1.22E+01)	$4.0002E {+} 02 (1.39E {-} 01)^{\dagger}$	4.7118E+02(1.69E+01)	$4.0000E+02(0.00E+00)^{\dagger}$
$f_{12}$	avgEvpRank	5616	3098	4250	4152	5987 <sup>†</sup>
	performEval	$5.0530E+02(3.35E+00)^{\dagger}$	3.3316E+03(5.94E+02)	5.6013E+02(6.08E+01)	5.0315E+03(5.90E+02)	$5.0481E+02(2.04E+00)^{\dagger}$
$f_{13}$	avgEvpRank	$4924^{\dagger}$	3782	4400	2876	4381
	performEval	$6.0000E+02(1.27E-04)^{\dagger}$	6.0003E+02(1.41E-02)	6.0000E+02(3.94E-04)	6.0161E+02(4.00E-01)	6.0002E+02(3.10E-03)
$f_{radar}$	avgEvpRank	2394	3601 <sup>†</sup>	1528	1316	1580
			$1.2365E+00(1.61E-01)^{\dagger}$	1.2614E+00(1.64E-01) <sup>†</sup>	1.9877E+00(1.55E-01)	1.9024E+00(8.60E-02)

The selected algorithms are highlighted in boldface.

# C. Performing the Algorithm Selection Task via the Performance Evaluation Approach

The algorithm selection task can also be performed by evaluating the performances of all candidate algorithms on a given optimization problem. In this paper, 50 independent runs of each candidate algorithm on each test problem were executed with a preset FEs budget of  $10^4 \times D$ . Then, statistical tests similar to those described in Section V-B3 were applied to compare the performances of the candidate algorithms on that problem. Concretely, the Kruskal-Wallis test was first used to check for differences among the entire set of candidate algorithms. If the null hypothesis of the Kruskal-Wallis test was rejected based on the preset significance level, the Conover-Iman test was further employed as a post hoc procedure for pair-wise comparisons of stochastic dominance. Then, the Holm procedure was applied to adjust the p-value obtained from the Conover-Iman test. Finally, the best algorithm or a set of equally best algorithms was selected based on the final results of these statistical tests.

# D. Results for the Algorithm Selection Task

For each test problem, the best algorithms were selected based on the proposed algorithm selection framework and the performance evaluation approach. As suggested

by Naudts and Kallel [53], the results obtained through performance evaluation can be regarded as the gold standard for the algorithm selection task, against which we can evaluate the results obtained by the proposed framework. As both approaches are based on statistical analysis, the evaluation criterion is as follows: for a test problem, if one of the best algorithms selected by the proposed framework is contained in the algorithm set chosen based on the performance evaluation approach, we consider the former to be correct; otherwise, it is considered a mistake.

Table II shows the selected algorithms for each test problem based on the two previously described approaches. For the proposed framework, the average rank in each sequence (i.e.,  $\overline{R_i}$ ) is shown as the statistical indicator. For the performance evaluation approach, we show the mean and standard deviation of the best fitness values obtained in 50 independent runs. As the statistical analysis procedures used for both approaches are the same, the results are comparable in terms of statistical significance. According to the evaluation criterion described above, the proposed approach achieves remarkable performance (with an accuracy of 100%) in the algorithm selection task. Moreover, in this paper, the FEs budget for online sampling and the number of neighbors for calculating the evp values were set to samFEs = 20%maxFEs and M = 10, respectively, and consequently, the proposed

<sup>&</sup>quot;avgEvpRank" and "performEval" denote the average evp rank in each sequence (i.e.,  $\overline{R_i}$ ) and the performance evaluation approach, respectively.

In all "performEval" rows, every measure is presented as the mean and standard deviation (in parentheses) for 50 independent runs.

<sup>&</sup>quot;in indicates that the selected algorithms are statistically significantly superior to the unselected algorithms at the 0.05 significance level and the 0.05 FWER level.

TABLE III
RELATION BETWEEN THE RESULTS OF THE ALGORITHM SELECTION TASK AND THE NUMBER OF NEIGHBORS

M	20	15	10	5	4	3	2	1
MisFunc	/	/	/	/	$f_7, f_{10}$	$f_7, f_{10}, f_{11}$	$f_4, f_7, f_9, f_{10}, f_{11}$	$f_4, f_5, f_7, f_8, \\ f_9, f_{10}, f_{11}, f_{radar}$
Cnt	0	0	0	0	2	3	5	8

<sup>&</sup>quot;M" denotes the number of neighbours.

TABLE IV
RELATION BETWEEN THE RESULTS OF THE ALGORITHM SELECTION TASK AND THE ONLINE SAMPLING BUDGET

samFEs	10%maxFEs	20%maxFEs	30%maxFEs	40%maxFEs	50%maxFEs
MisFunc	$f_8, f_{10}, f_{13}$	/	/	/	/
Cnt	3	0	0	0	0

<sup>&</sup>quot;samFEs" denotes the budget of FEs for online sampling.

algorithm selection framework incurred a much lower computational cost than the performance evaluation approach. We will discuss the details of further reducing the computational cost in the next section.

The detailed differences between the algorithms selected using the two approaches can also be explored based on Table II. Concretely, for the unimodal test problems, i.e.,  $f_1$ – $f_5$ , the proposed framework selects only one algorithm, whereas more than two algorithms are selected by the performance evaluation approach. After further checking the number of FEs required for the convergence of algorithms, we discover that the proposed framework prioritizes the algorithm that has the best performance and also converges fastest. This phenomenon can be easily understood, as the best-performing and fastest-converging algorithm has the highest evp values. For the multimodal test problems, i.e.,  $f_6$ – $f_{13}$ , the proposed framework always selects the best-performing algorithm in terms of the best mean and the lowest standard deviation. Although other algorithms, such as CoDE, are also selected by the performance evaluation approach for several problems, the proposed framework tends to select the best-performing (i.e., with the best mean) and the most robust (i.e., with the lowest standard deviation) algorithm. Notably, the proposed framework can make the correct choice even on the NP-hard numerical optimization problem (i.e.,  $f_{radar}$ ), which indicates its general applicability for the algorithm selection task.

# VI. FURTHER DISCUSSION

# A. Robustness of the Proposed Framework

Here, the robustness of the proposed framework for the algorithm selection task is discussed. As mentioned earlier, the proposed framework for the algorithm selection task is based on online sampling and estimates of the evp measure. Both online sampling and measure estimation procedures contain randomness, and seed of the random number generator is reset based on the current time before either procedure is performed. Therefore, if the results are robust to this intrinsic randomness, the proposed algorithm selection framework can be considered reasonable and trustworthy.

An experiment was conducted to test the robustness of the proposed framework. In this experiment, we first fixed the FEs budget for online sampling and the number of neighbors to be generated for estimating the evp values to samFEs = 20%maxFEs and M = 10, respectively, and then ran the proposed framework 10 times. The results obtained from each run were statistically evaluated with respect to those achieved using the performance evaluation approach (shown in Table II). We found that the results obtained using the proposed framework are stable. Specifically, the proposed framework achieved an accuracy of 100% three times and made one mistake four times. For the remainder of the runs, it made more than one but fewer than four mistakes, most of which occurred for the functions  $f_8$ ,  $f_{10}$ , and  $f_{13}$ . Notably, these functions that confuse the proposed framework are all complicated multimodal functions with many local optima, which may affect the information obtained via online sampling. However, in all of the cases in which the proposed framework failed to choose the correct algorithm, it ranked that algorithm second, i.e., was not far off.

# B. Accuracy of the Proposed Framework Versus Computational Cost

In this section, two tunable parameters of the proposed framework are discussed. The first one is the number of neighbors (i.e., M) used for measure estimation. This parameter can be interpreted as specifying the tradeoff between the accuracy of estimation and the computational cost: the larger the number of neighbors is, the more accurate the measure estimates should become, at the price of a higher computational cost. To further investigate the influence of this parameter on the accuracy of the framework for the algorithm selection task, we first fixed the FEs budget for online sampling to samFEs = 20%maxFEs and observed the relation between the results of the algorithm selection task and the number of neighbors.

As shown in Table III, if the number of neighbors M decreases below 5, the number of test functions for which the proposed framework makes mistakes will start to increase.

<sup>&</sup>quot;MisFunc" denotes the test functions on which the proposed framework made mistakes

<sup>&</sup>quot;Cnt" denotes the number of functions in MisFunc.

<sup>&</sup>quot;MisFunc" denotes the test functions on which the proposed framework made mistakes.

<sup>&</sup>quot;Cnt" denotes the number of functions in MisFunc.

Therefore, the neighbor number threshold that ensures the accuracy of the proposed framework is 5. More importantly, when samFEs and M are separately set to 20%maxFEs and 5, the total number of FEs required to execute the entirety of the proposed framework for the algorithm selection task is equal to or fewer than maxFEs, which is at most sufficient to perform one run in the performance evaluation approach. Generally, performing only one run for each metaheuristic algorithm in a black-box optimization problem is not sufficient for a statistically sound determination of which algorithm is the best choice for that problem, whereas the proposed framework can produce statistically significant results with high accuracy at a similar or lower computational cost in terms of FEs.

For the other parameter, i.e., the FEs budget for online sampling (samFEs), a similar study was performed with the number of neighbors set to M=10. As illustrated in Table IV, when samFEs is set to 10%maxFEs, the proposed approach makes three mistakes, for functions  $f_8$ ,  $f_{10}$ , and  $f_{13}$ . When samFEs is increased from 20%maxFEs up to 50%maxFEs, the proposed framework remains an accuracy of 100%. It should be noted that additional experiments with continuously increasing values of this parameter were not performed because such values would markedly increase the computational cost of the proposed framework. In this paper, we recommend samFEs be set to 20%maxFEs.

#### VII. CONCLUSION

In this paper, the concept of population evolvability is presented as an extension of dynamic FLA. Two measures, i.e., epp and eap, are proposed to describe the probability that a population will obtain improved solutions to an optimization problem and its ability to do so. Then, a combined measure, i.e., evp, is derived to represent the overall population evolvability. Experimental studies illustrate the validity of the proposed measures. The algorithm selection task for black-box optimization problems is considered as an application to demonstrate that the proposed framework is a generally applicable and low-cost approach for performing the algorithm selection task with high accuracy and statistical significance.

It is noteworthy that this paper is merely a preliminary step toward the goal of developing more practical FLA techniques. Additional studies based on the proposed concept will be conducted in the future. From the perspective of algorithm designers, further work can be focused on the design and application of the proposed measures for the hyper-parameter tuning of population-based metaheuristic algorithms, including parameter control, algorithm portfolios and hybridization.

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