

# A NEW APPROACH TO DETECTING AND PREVENTING POPULATIONS STAGNATION THROUGH DYNAMIC CHANGES IN MULTI-POPULATION-BASED ALGORITHMS

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#### Abstract

In this paper, a new mechanism for detecting population stagnation based on the analysis of the local improvement of the evaluation function and the infinite impulse response filter is proposed. The purpose of this mechanism is to improve the population stagnation detection capability for various optimization scenarios, and thus to improve multi-population-based algorithms (MPBAs) performance. In addition, various other approaches have been proposed to eliminate stagnation, including approaches aimed at both improving performance and reducing the complexity of the algorithms. The developed methods were tested, among the others, for various migration topologies and various MP-BAs, including the MNIA algorithm, which allows the use of many different base algorithms and thus eliminates the need to select the population-based algorithm for a given simulation problem. The simulations were performed for typical benchmark functions and control problems. The obtained results confirm the validity of the developed method. Keywords: multi-population-based algorithms, migration topologies, population stagnation

# 1 Introduction

Population-based algorithms allow for iterative processing of a population of solutions (called individuals) to find the most optimal solution. The

solutions are encoded by the parameters of the individuals, while their quality is determined by the evaluation function associated with the problem under consideration. The parameters encoded by indi-

viduals are modified by various mechanisms (operators) that can be local (exploitation), global (exploration), or everything in between. In modifying the parameters of one individual, only random or auxiliary variables (e.g. mutation or momentum, respectively) or parameters of other individuals (e.g. crossover) may be used. The use of parameters of other individuals, or the use of small changes in parameters, can cause the whole population to become very similar in terms of these parameters, and the solutions converge to one area of the parameter space (search space). Thus populations can often stuck in the local optimum. Several mechanisms can be used to prevent this, and one of them is the simultaneous use of multiple populations, the approach considered in this paper.

Multi-population-based algorithms (MPBAs) use multiple populations (also called islands or subpopulations) that are usually processed independently, while in certain situations individuals between them are exchanged (migrated). The methods of sub-populations processing are determined by the training plans. The selection of the populations for the exchange of individuals is determined by the topology of their connections (migration topology). The number of exchanged individuals and the method of their selection is determined by the migration strategy, while the replacement period is determined by the frequency of migration. Thus, many factors can affect the performance of multi-population-based algorithms (including selecting basic optimization parameters), which makes the problem of their configuration complicated and may lead to premature stagnation. The more details on the problems associated with MPBAs are described in the next section. In this paper, an attempt was made to further improve MP-BAs in various aspects, proposing more universal solutions, and including using mechanisms that try to prevent premature convergence.

#### 1.1 Motivation

The high popularity of population-based and multi-population-based algorithms has its advantages and disadvantages. On the one hand, new and better algorithms appear frequently, and algorithms are adapted to more and more complex simulation problems (see e.g. [14]). On the other hand, the multitude of algorithms and their versions makes

it difficult to choose the best algorithm for a given simulation problem (see e.g. [7, 15, 18, 43]). A large number of population-based algorithms also makes it difficult to compare them, and most often in the papers on a new version of an algorithm, it is compared only with related algorithms (or their modifications) or original versions of other algorithms (see e.g. [24, 37]). This also applies to various mechanisms used in multi-populationbased algorithms, such as the development of new topologies and migration strategies, mechanisms to prevent stagnation, etc., which are sometimes only tested for one population-based algorithm or tested for multiple algorithms but not compared with other strategies. Such papers, however, then focus on other aspects of the operation of the considered algorithms (see e.g. [1, 17]). Of course, these problems arise mainly due to the difficulty of running such complex simulations and keeping track of all the latest versions of various population-based algorithms, and thus they are practically inevitable.

In this paper, it was decided to focus on universality. Firstly, the MNIA algorithm is further developed, in which many sub-populations based on different population-based algorithms can cooperate. Secondly, new approaches were proposed for stagnation detection and the dynamic changes in active populations and the number of their individuals. These solutions are designed to apply to most migration topologies and various MPBAs. Thirdly, the proposed approaches, in addition to a comparison with known mechanisms, will be tested for various migration topologies and various sub-population algorithms. Such assumptions, besides the proposal of new mechanisms, should not only find more efficient configurations of algorithms that give better optimization results but also determine the further possible direction of development of such algorithms.

### 1.2 Contribution

The contribution of this paper to the development of population-based algorithms is as follows: a) directing further development of more universal methods that allow to eliminate the problem of selecting a specific population-based algorithm, in particular, the development of the MNIA algorithm [40]; b) proposing and applying known and new mechanisms related to the dynamic change in the number of populations and their individuals, aimed at improving the accuracy of population-based algorithms; c) proposing and applying known and new mechanisms related to the dynamic change in the number of populations and their individuals, aimed at improving the complexity of populationbased algorithms; d) development and testing of a new criterion to determine the level of population stagnation, in particular for use in a multipopulation-based algorithms; e) a detailed and comprehensive comparison of different methods for changing the number of populations and the number of individuals used with different migration topologies and for different multi-population-based algorithms.

#### 1.3 Paper Structure

The structure of the paper is as follows: Section 2 presents a review of the literature, Section 3 describes the proposed method, Section 4 presents considered simulation variants and obtained results, and Section 5 draws final conclusions.

### 2 Background

According to the no-free lunch theory [2], there is no single population-based algorithm that is best suited for all simulation problems. This, on the one hand, causes a problem because a lot of new population-based algorithms are being developed, which is already criticized (see e.g. [7, 15, 43]). An overview of various algorithms can be found e.g. in [15, 23, 38, 45]. On the other hand, new algorithms are developed in such a way as to improve the search mechanisms and avoid premature convergence of the algorithm (stagnation of the population at the local optimum), and therefore this is the positive part. New algorithms have more configuration options, although there are exceptions, e.g. [31]). Moreover, new algorithms can be more computationally complex, which can be critical when optimizing more complicated simulation problems (e.g. control systems problems [25, 32], fuzzy systems [41, 47]). Moreover, the spectrum of applications of these algorithms is constantly increasing (see e.g. [48]). At the same time, some algorithms combine the advantages of different algorithms and become universal, partially eliminating the problem of choosing a specific population-based algorithm

for a specific simulation problem (see e.g. [40]). Despite the rapid development of population-based algorithms, the problem of premature convergence (stagnation) still occurs and is crucial.

One way to prevent population stagnation in population-based algorithms is to diversify the behavior of the population. In the simplest approaches with one population individuals can be divided according to certain criteria. Examples are Grey Wolf Optimizer, in which three main individuals and the rest of the herd are distinguished [31], Bison Algorithm, in which there are individuals with two different behaviors [21], or Termite Queen Algorithm, in which there are as many as five different types of individuals (queen, flying worker, foraging worker, serving worker, and soldier [9]). In some cases, distinguished individuals can be modified using different methods of changing parameters that originate from different population-based algorithms, and thus hybridization occurs (see e.g. [11, 14, 27]).

The advanced approach consists of dividing the population into separated sub-populations (islands) that exchange solutions in certain circumstances. Usually, islands operate based on the same population-based algorithm, and the method of modifying individuals on each island may be identical (see e.g. [4]), or depending on the parameters of a given island (see e.g. [36]). There are also solutions where the behavior of each island can be based on different operators or different population algorithms (see e.g. [40]). When only two or three sub-populations are used and a different populationbased algorithm is used for each of them, such solutions are also called hybrid (see e.g. [33]). In the case of a larger number of sub-populations, it becomes crucial to select the topology of their connections determining the way of exchanging individuals and other mechanisms described later in this section, broken down into topologies, migration strategies, parameters, adaptive mechanisms, and hybridizations with other, non-population-based, algorithms. Each of these mechanisms can have a significant impact on the algorithm's effectiveness and thus is important from the point of view of this paper.

### 2.1 Migration topologies

There are many ways to improve the performance of multi-population-based algorithms. The first one is the selection of the appropriate migration topology. Different topologies have different effects on population diversification and search intensification. The selection of the appropriate topology can affect the balance between exploration and exploitation and also affect it differently depending on the simulation problem under consideration.



Figure 1. Efficient migration topologies according to [12]: a) inverted star with the best population in the middle (ISBM), b) ladder with a down-top connection (LDTC), c) topology with two random connections for each population (TTRC), d) inverted star as in a) combined with ring (IRBM) e) ring with additional connections between neighbors (RWAC), f) ring with additional random connections, also called small world (SMWD). Where: rectangles stand for populations, dark rectangles stand for populations that are conditionally replaced with best population, arrows stand for migration directions, and dotted arrows stand for random connections.

The most common topologies according to [19] are star and ring, while others worth noting are torus and lattice. Some topologies also change dynamically, examples are randomly changed topologies (see e.g. [8]) or topologies where the best or worst population is placed in a specific location (see e.g. [12]). Moreover, different approaches, such as a

fitness-based Migration Policy designed to promote the maintenance of diversity through a mechanism that combines groups of individuals to alternate between exploration and exploitation proposed in [6], can be used instead.

An interesting overview and performance tests of 36 different migration topologies is presented in [12]. The best topologies, regardless of other configurations and simulation problems used, turned out to be: an inverted star with a mechanism for setting the best population in the middle of topology (see Figure 1.a)), a ladder with a down-top connection (see Figure 1.b)), and a topology in which each population has two random connections with others (see Figure 1.c)). Due to their universality, these topologies will be used in the simulations in this paper.

### 2.2 Migration strategies

Sub-populations migration is a key element of island algorithms. This mechanism may relate to the frequency of migration, the number of individuals to be exchanged, and the method of selection of individuals for exchange. Not only does population diversification depend on these configurations, but also the accuracy of the search. In addition to the standard configuration described above and specifying methods for selecting and replacing individuals, other solutions are also used. In [35] the individuals of the two populations are mixed randomly at each iteration of the algorithm, and in [13] individuals are moved into a dynamic number of sub-populations instead. In [49] a grouping mechanism is used to divide the population into two subswarms. In the [30] the authors use a superior population from which they send non-dominated solutions to subpopulations.

The paper [12] shows the influence of migration parameters on a given migration topology. The best parameters turned out to be the frequency of replacement every 10 iterations of the algorithm, and the selection of an individual for replacement using the roulette wheel method. With such parameters, the following topologies turned out to be worth nothing: an inverted star with a ring and placing the best population in the middle (see Figure 1.d)), a ring with additional connections between neighboring populations (see Figure 1.e)) and a ring with two additional random connections (see Figure 1.f)).

These topologies were also considered in this paper's simulations.

### 2.3 Parameter adjustment

The selection of parameters for MPBAs is the third important factor in fine-tuning these algorithms. This includes, among others, the selection of the number of islands and the number of individuals in these islands. Choosing the number of individuals is not an easy task, because each population-based algorithm works differently for different numbers of individuals (some algorithms need fewer individuals and more iterations, and vice versa). The parameters of specific islands are also important, which may be the same for each island or different. What is more, the operation of each island can be based on mechanisms from other populationbased algorithms.

Parameter tuning can be done offline (selection of parameters before applying the algorithm) or online where parameters are changed or adapted during the simulations [16]. In the case of offline parameter selection, they can be set by trial and error to static values, or values depending on a given step of the algorithm (see e.g. [11]). In the case of MPBAs, different parameters can be set for each population and thus increase the chance of adapting the algorithm to a given problem (see e.g. [40]). Another parameter optimization idea is metaoptimization, which is the optimization of parameters for some underlying optimizer using a different algorithm [16, 26]. This approach is also used in multi-criteria algorithms, where, for example, algorithm parameters are optimized in such a way as to obtain the best results of multi-criteria performance metrics [26].

The paper [12] shows that, regardless of the migration topology and the simulation problem under consideration used, a good configuration is a topology with 8 islands composed of 32 individuals each, while the paper [40] shows that algorithms in which islands operate based on different population algorithms allow obtaining better and more stable results.

### 2.4 Adaptive mechanisms

Adaptive mechanisms mean dynamic adjustment of the algorithm operation through all configurations (migration topology, migration strategy, and algorithm parameters). Modification of the above settings is most often done based on the detection of population stagnation and optimization progress analysis. Some of these mechanisms are also used for common single-population-based algorithms.

In [46] Differential Evolution parameters are adaptively adjusted according to the statistical information learned from the previous searches in generating improved solutions. In [5] fuzzy system is used to control population diversity at decision variable space by self-adapting the crossover rate control parameter. In [22] an adapted crossover rate value is assigned to each individual according to individual fitness value. It is also worth mentioning the adaptive mechanisms of re-initializing the population (or part of it), which allow one to search subsequent areas of search space and thus to improve the results (see e.g. [29]). Adaptive mechanisms can be focused also on the complexity of the optimization - e.g. by dynamically changing the number of populations or individuals, as well as a dynamic selection of surrogate solutions (see e.g.  $[11]$ ).

In the [28], individual island training plans (both population algorithms and their parameters) are changed based on the optimization progress, which brought a significant improvement. In the paper [29], when stagnation was detected, various mechanisms were used to re-initialize those populations that performed the worst. In this case, the best solution turned out to be re-initialization involving the creation of new individuals based on the mechanisms of differential evolution and the use for re-initialization the individuals from other populations. In this paper, it is planned to test the use of different approaches to adaptive changes in the MP-BAs.

### 2.5 Hybridizations

In addition to the aforementioned hybridization consisting of the simultaneous use of different population-based algorithms or operators derived from them, it is worth mentioning that the literature is full of methods in which populationbased algorithms are combined with other families of algorithms. Hybridizations worth mentioning are the combination of population-based algorithms with memetic algorithms, which are also evolutionary algorithms but focused on local search (see e.g. [34, 44]) or algorithms based on backpropagation (see e.g. [3, 20]). This shows how important the population-based methods are and that there is still a huge potential for their development. Such approaches will not be considered in this paper due to their complexity, however, it is possible to explore such an interesting topic in the future.

## 3 Proposed approach

Section 2 shows how many different approaches can be used to improve the performance of MPBAs, how many parameters influence the performance of these algorithms, and what approaches allow them to obtain promising results. This section describes the proposed approach and its configuration with its division into the mechanisms of changing the number of populations and the number of individuals within the populations, mechanisms of preventing population stagnation, and a complete description of the proposed algorithm. The proposed approach has been developed in such a way that it is as universal as possible and can be used for many topologies, algorithms, and problems.

### 3.1 Dynamic number of populations and individuals within populations

There are many approaches for changing population and individual numbers (see also Section 2.2). The purpose of such changes may be to increase the number of individuals in more promising populations, remove less promising populations or add new populations, increase the diversity of individuals, or reduce computational complexity, all to improve the performance of MPBAs. Some methods are typical for specific algorithms and do not allow their use in different topologies. For example, the ICA [4] algorithm has individuals grouped into empires with an imposed migration topology, in which the best population gains an individual from the worst population, thus the number of groups in the population may slowly decrease and thus it is not possible to apply other topologies directly.

In this paper, the use of the following universal approaches is proposed allowing for their use in most migration topologies: a) disabling or enabling of specified populations - in this case, the disabled populations are not iterated, but to ensure compliance with the migration topology, they may continue to participate in the transfer of individuals, b) dynamic changes in the number of individuals of selected populations - if there is a need to reduce the number of individuals, the worst solutions in terms of adopted evaluation function are removed, and if there is a need to increase the number of individuals, new ones are created based on tournament selection of individuals from other populations and the use of a mutation operator derived from the genetic algorithm (to maintain population diversity). The above mechanisms can be used in any combination (see Table 1) and when certain conditions occur (e.g. stagnation detection). The idea of their use is as follows: populations are sorted according to the evaluation function of the best individuals, the number of population individuals is checked against the number of individuals resulting from the adopted strategy (see Table 1), if individuals need to be added, removed or a given population needs to be disabled or enabled, the approaches described above are used.

The proposed approaches were divided into two categories: focused on improving accuracy (IA) and focused on reducing complexity (RC). Their combinations are presented in Table 1. The first approach involved increasing individuals in *Nchg* populations while extinguishing individuals in other *Nchg* populations (for the details see DBDSW and DSDSB in Table 1). The next goal is just to extinguish individuals in *Nchg* populations and thus reduce complexity (see DSBST and DSWRS in Table 1). Then, the use of a solution was proposed in which there are no disabled populations but only changes in the number of individuals - increasing the number of individuals in the 2*·Nchg* populations and reducing them in the second 2*·Nchg* populations (see LWIBS and LBIWR in Table 1). This variant was also presented similarly in the version with complexity reduction (see LWRWR and LWRBS in Table 1). The latest variants were designed with a more linear change in the number of individuals in the analogous matter (see LDFBW, LDFWB, LDRBW, and LDRWB in Table 1). The above changes were designed so that the number of individuals in the IA approach is constant, while in the RC approach, it was reduced by 25% compared to the variant aimed at improving accuracy. It is worth noting, that despite assumptions about how to add or remove indi-

Table 1. Mechanisms for changing the number of populations and individuals within populations used to improve accuracy or complexity (see Section 3.1). Examples are given for *N pop*=8*,Nind* =32*,Nchg*=2.

strategy	aim	description	examples of population sizes ordered from best to worst	processed individuals
<b>DBDSW</b>	<b>IA</b>	disable <i>Nchg</i> best populations, double size of <i>Nchg</i> worst populations	$-$ , $-$ , 32, 32, 32, 32, 64, 64	100%
<b>DWDSB</b>	IA	disable <i>Nchg</i> worst populations, double size of <i>Nchg</i> best populations	64, 64, 32, 32, 32, 32, -, -	100%
<b>DSBST</b>	<b>RC</b>	disable Nchg best populations	$-$ , $-$ , 32, 32, 32, 32, 32, 32	75%
<b>DSWRS</b>	RC	disable Nchg worst populations	$32, 32, 32, 32, 32, 32, -$	75%
<b>LWIBS</b>	<b>IA</b>	lower $2 \cdot Nchg$ worst, increase size of $2 \cdot Nchg$ best	48, 48, 48, 48, 16, 16, 16, 16	100%
<b>LBIWR</b>	<b>IA</b>	lower $2 \cdot Nchg$ worst, increase size of $2 \cdot Nchg$ best	16, 16, 16, 16, 48, 48, 48, 48	100%
<b>LWRWR</b>	RC	lower $2 \cdot Nchg$ worst individuals	32, 32, 32, 32, 16, 16, 16, 16	75%
<b>LWRBS</b>	RC	lower $2 \cdot Nchg$ best individuals	16, 16, 16, 16, 32, 32, 32, 32	75%
<b>LDFBW</b>	IA	linear individuals distribution, from best to worst	60, 52, 44, 36, 28, 20, 12, 4	100%
<b>LDFWB</b>	<b>IA</b>	linear individuals distribution, from worst to best	4, 12, 20, 28, 36, 44, 52, 60	100%
<b>LDRBW</b>	RC	half-linear distribution with reduction, from best to worst	32, 32, 32, 32, 28, 20, 12, 4	75%
<b>LDRWB</b>	<b>RC</b>	half-linear distribution with reduction, from worst to best	4, 12, 20, 28, 32, 32, 32, 32	75%

viduals, the proposed general approach does not determine a rigid definition of these mechanisms and any other approaches can be used instead.

### 3.2 Prevention of population stagnation

The mechanisms presented in Section 3.1 can be used in various situations. In this paper, it was assumed that their main purpose was to improve MPBAs and to prevent stagnation. Therefore, it is crucial to determine at what point in the optimization they should be used.

In the basic approach (STA), dynamic change mechanisms are triggered after a certain number of iterations of the algorithm. In more advanced approaches, various mechanisms for detecting stagnation can be used. One of them is detecting whether there has been an improvement in the evaluation function over the *Nimp* iterations (SFF). In addition to detecting stagnation, it is also possible to use population convergence techniques that are based on the similarity of individuals. However, all of the mentioned mechanisms may be disturbed, because even a small improvement in accuracy results in the lack of use of given mechanisms. The solution is to check the improvement above a certain threshold or to calculate the average fitness of the population, thus the problem of determining the threshold for a given simulation problem and the problem of disturbance of the average fitness by migrations arise, respectively. In addition, in the case of multi-population-based algorithms, it is necessary to choose whether these mechanisms should be used separately for each population or whether to test the general stagnation of all populations. In the first approach, it is not possible to use some approaches to prevent stagnation (e.g. in the case of a change in the number of individuals of all populations), thus it is not used in this paper.

To prevent the above problems, a new criterion for stagnation detection, resistant to both minor and abrupt changes in the improvement of the evaluation function, was proposed in this paper. The use of standard improvement is difficult due to, for example, the possibility of the fitness value falling below zero (thus logarithmic approaches do not always apply) as well as the aforementioned problem of analysis of a given simulation benchmark (it is required that it works equally well at the beginning of the simulations when the changes in the fitness

function values are large, as well as in the final stage of optimization, and regardless of the range of values of a given simulation problem). To solve that, first, a flexible percentage improvement in fitness values between iterations is used as a basis. The proposed flexible percentage is calculated for *Nels* of the last iterations of each population and checks the level of improvement of the evaluation function relative to the improvement from *Nels* optimization steps:

$$
imp_i(iter) = \frac{ff_i^{bst}(iter-1) - ff_i^{bst}(iter)}{ff_i^{bst}(iter-Nels) - ff_i^{bst}(iter)},
$$
 (1)

where  $\text{ff}^{bst}_i(iter)$  is the best fitness function value of individual from *i*-th population for *iter* iteration step  $(i = 1, ..., Npop, Npop$  is the number of subpopulations), *Nels* is the adopted time window. It is worth noting that in the case of the above equa- $\text{tion, if } \text{ff}^{bst}_i(iter - stp) - \text{ff}^{bst}_i(iter) = 0, \text{ the result}$ should be equal to zero, and if  $iter < stp$ , the default value of the improvement γ should be used. To immunize the above improvement factor to small and large fluctuations in the improvement of the evaluation function, an infinite impulse response filter was used (see [10]). In this case, the general formula for proposed stagnation detection for single population is defined as follows:

$$
stgi(iter) = \alpha \cdot stgi(iter-1) + (1-\alpha) \cdot stgi(iter), (2)
$$

where  $\alpha$  is a coefficient determining how much the value from the previous iteration step has an impact on the proposed criterion. In the case of a first iteration step  $stg_i(1)$ , the value is set to the coefficient γ value. The stagnation in the proposed approach can be detected by checking if the  $stg_i(tier)$  value has dropped below a certain β value. Examples of the calculated ratio are shown in Figure 2. As can be seen in Figure 2, the proposed value  $stg_i(tier)$ works well in both presented cases. It is worth noting that in case Figure 2.a) there is often a small improvement in the population, making the SFF approach ineffective and detects stagnation only towards the end, while in case of Figure 2.b) there is more downtime during optimization, sometimes after a big improvement - in this case, using the standard approaches may be too frequent. The proposed approach (ELA) is new in the literature, in partic-



Figure 2. Example of population stagnation detection: area marked in a dark color - standard approach, area marked in a light color - proposed approach, *stg* stands for proposed criterion (see equation (2)), and *nni* stands for the number of iterations without improvement: a) slow and steady improvement in the middle of the optimization, b) more dynamic but stable changes in the fitness values.

ular in terms of performance evaluation of multipopulation algorithms.

### 3.3 Proposed algorithm

The approach proposed in this paper has been designed universally so that most migration topologies, stagnation repair mechanisms, or populationbased algorithms can be used.

The algorithm consists of the following steps:

Step 1. Initialization of *N pop* populations and for each of them an equal number of *Nind* individuals. Both the initialization method and the choice of population processing algorithms can be completely arbitrary. In the case of the MNIA approach, each population can be processed by a different algorithm with different parameters. In this step, individuals are also evaluated and the base number of individuals and population status (disabled/enabled) are changed according to the rules defined by Table 1 and the optimization parameters. After initialization, the populations are set up according to the adopted migration topology (see Fig 1). Setting the *iter* variable to 0.

Step 2. In this step, the processing of individuals of each population is triggered by the adopted population-based algorithm. This step can be processed a certain number of iterations (*Niter*) or until a certain optimization fitness function value error threshold is reached. At the end of a single iteration, the *iter* variable is incremented and the following conditions are checked:

- Checking whether the migration topology needs to be changed, e.g. in the case of the ISBM topology, the best solutions are transferred to the central point of the topology. Changes are checked and applied in each iteration of the algorithm.
- Checking if *Nmig* iterations have elapsed. If so, the exchange of individuals between populations is by the adopted topology and migration strategy.
- Checking the conditions determining the need to change the number of active populations or the number of individuals within them according to the proposed mechanism of reducing and adding individuals (see Section 3.1). If the conditions are met for at least *Nwar* of the sub-populations, the strategies presented in Table 1 apply. The examples of such changes are shown in Table 2.

Table 2. An example of changes in populations after stagnation is detected and the populations are sorted according to the evaluation function of their best individuals for the DWDSB case,  $Npop = 8$ ,  $Nind = 32$ 

and *Nchg* = 2 (see Table 1). The number of individuals and population statuses results from previous changes where the populations may have been sorted in a different order, '-' stands for disabled population.



Step 3. Checking if the stop condition is reached, if not return to step 2. If so, present the best-found solution in terms of the adapted fitness function.

The algorithm considered in this paper does not require a specific approach to encoding solutions and their evaluation and these issues will not be considered in detail. However, the algorithm can be easily adapted, to solving hybrid-type problems (consisting of finding the solution structure and its parameters), etc. Then, the method of encoding solutions may be analogous to that presented in the papers [25, 40].

# 4 Simulations

The simulations were carried out in such a way as to test both the strategies from Table 1 and compare different approaches to preventing stagnation (including the proposed one), all for different migration topologies, different population algorithms, and various simulation problems.

The assumptions and parameters of the simulation are as follows: topologies tested: ISBM, LDTC, TTRC, IRBM, RWAC, SMWD (see Figure 1), all strategies tested from Table 1, for the migrations one worst individual from one population is replaced by one individual selected by roulette wheel from other randomly selected population. Used multi-population-based algorithms: island algorithms with same population-based in all populations MPGA, MPDE, MPCS, MPGWO (with consecutively base algorithms: GA, DE, CS, GWO with default parameters from the literature), MNIA (GA+DE+CS+GWO). Base number of populations

being changed after stagnation detection *Nchg* = 2, number of iterations for STA approach *Nsta* = 25, number of iterations without improvements for SFF approach *Nimp* = 6, parameters for proposed method ELA:  $\alpha$  = 0.9, *Nst p* = 10, β = 0.06, γ = 0.2, number of populations with stagnation to apply prevention mechanisms *Nwar* = 2, number of islands  $Npop = 8$ , based number of individuals in islands *Nind* = 32, number of iterations *Niter* = 1000, number of simulation repetitions *Nrep* = 50, migration interval *Nmig* = 10. The mentioned parameters of the above approaches were determined by trial and error method (see also Figure 3) or based on the values suggested in the literature, however, in the future, it is possible to test the meta-optimization approach, in which the parameters will be selected automatically.



Figure 3. Influence of parameters of ELA and SFF approaches on fitness function values improvement

(ffimp). In the case of a), b) and c), parameters other than those tested were set as follows:  $\alpha = 0.9$ ,  $β = 0.06, γ = 0.2.$ 

Table 3. Results averaged across all benchmark functions and all multi-population-based algorithms. The results show the improvement in results (in percent) relative to the same methods without the use of stagnation detection and prevention mechanisms. SM stands for stagnation mechanisms, SS stands for stagnation strategy. The best results for the accuracy approach are in bold, and for the complexity reduction approach are underlined.

<b>SM</b>	<b>SS</b>	<b>ISBM</b>	<b>LDTC</b>	<b>TTRC</b>	<b>IRBM</b>	<b>RWAC</b>	<b>SMWD</b>	<b>AVG</b>
<b>STA</b>	<b>DBDSW</b>	147	173	143	243	254	199	193
	<b>DWDSB</b>	216	118	120	150	$-154$	69	86
	<b>DSBST</b>	33	41	50	50	52	47	46
	<b>DSWRS</b>	52	49	62	77	59	85	64
	<b>LWIBS</b>	185	148	139	222	201	244	190
	<b>LBIWR</b>	773	337	282	$-111$	$-94$	39	204
	<b>LWRWR</b>	111	96	94	141	123	129	116
	<b>LWRBS</b>	160	129	90	154	142	147	137
	<b>LDFBW</b>	74	173	150	252	281	181	185
	<b>LDFWB</b>	259	233	241	407	640	338	353
	<b>LDRBW</b>	122	113	90	162	124	125	123
	<b>LDRWB</b>	132	107	117	152	134	113	126
	<b>DBDSW</b>	118	$\overline{140}$	130	147	145	144	137
	<b>DWDSB</b>	86	69	89	96	87	88	86
	<b>DSBST</b>	46	42	47	51	57	89	55
	<b>DSWRS</b>	118	57	75	64	66	53	72
	<b>LWIBS</b>	152	137	125	193	183	165	159
<b>SFF</b>	<b>LBIWR</b>	181	203	520	269	257	$-26$	234
	<b>LWRWR</b>	114	105	86	160	117	108	115
	<b>LWRBS</b>	107	115	100	180	156	149	134
	<b>LDFBW</b>	187	161	131	269	190	194	189
	<b>LDFWB</b>	173	288	$-579$	286	301	215	114
	<b>LDRBW</b>	111	114	86	146	132	114	117
	<b>LDRWB</b>	155	148	115	166	159	262	168
	<b>DBDSW</b>	649	261	142	709	207	250	370
	<b>DWDSB</b>	287	136	$-59$	259	147	111	147
	<b>DSBST</b>	34	33	129	46	58	49	58
	<b>DSWRS</b>	62	66	52	259	$70\,$	63	95
	<b>LWIBS</b>	204	155	152	121	220	199	175
<b>ELA</b>	<b>LBIWR</b>	201	589	313	32	165	238	256
	<b>LWRWR</b>	109	97	83	146	136	108	113
	<b>LWRBS</b>	121	138	113	165	146	145	138
	<b>LDFBW</b>	98	172	169	279	290	203	202
	<b>LDFWB</b>	312	664	362	$-188$	557	576	381
	<b>LDRBW</b>	135	120	95	153	145	128	130
	<b>LDRWB</b>	221	114	106	147	211	154	159
	<b>AVG</b>	173	162	116	168	166	153	156





Table 5. Results for the proposed stagnation method ELA averaged across all multi-population-based algorithms and stagnation strategies for the control problems. The results show the improvement in results (in percent) relative to the same methods without the use of stagnation detection and prevention mechanisms. SP stands for simulation problem. The best results are in bold.

SP.			ISBM LDTC TTRC IRBM RWAC SMWD AVG	

The considered methods were tested using the CEC2013 benchmark functions (improved CEC2005 benchmark set as used in [42]), hereinafter referred to as C01-C28 [39]. In addition, problems of selecting the parameters of control systems based on PID elements: MSD, WTT (see details in [25]) were also considered.

### 4.1 Simulation results

For a clearer presentation of the results and to show the capabilities of the mechanisms used, all results are shown as a percentage improvement over the case where no stagnation detection and prevention mechanisms were used. For example, a value of 100% means an improvement in error value from 1.0 to 0.5, a value of 200% from 1.0 to 0.25, and a value of -100% a deterioration from 1.0 to 2.0 (all problems concerned the minimization of the value of the evaluation function).

Results detailing all simulation cases are shown in Table 3, results detailing simulation problems for the proposed ELA stagnation detection method are shown in Table 4, results detailing various multipopulation-based algorithms are shown in Table 6, and general summary of the individual methods is presented in Table 7. In addition, Figure 3 presents the impact of the α, β, γ, and *Nimp* values on the results obtained for the ELA method and SFF method respectively. Results for control problems are shown in Table 5.

#### 4.2 Simulation conclusions

The best results in terms of fitness function values were achieved for the proposed stagnation detection strategy (ELA) and the LDFWB and DBDSW stagnation prevention mechanisms, while the proposed LDFWB mechanism also performed very well in the standard approach (STA) - see bold values in Table 3. For variants with less computational complexity, the best mechanism is LDRWB, but in this case, the stagnation detection approach based on the lack of improvement of the evaluation function (SFF) worked slightly better (see LDRWB on SFF and ELA in Table 3. For each migration topology a satisfactory improvement over the lack of stagnation detection mechanisms was achieved, the only exception is the TTRC topology - see Table 3.

According to the no-free lunch theory, different degrees of improvement were achieved for each simulation problem using the ELA mechanism and the stagnation prevention mechanisms. For some configurations, there was no improvement in the results (see negative values in Table 4). The ELA mechanism improved the performance of the optimization for each of the topologies considered, with the best topology (IRBM) improving by 177%, and the next topology (ISBM) improving by 203%, thus giving the best results overall - see Table 4. The ELA approach also improved the results for control problems (see Table 5). In the case of non-synthetic tests, the improvement was significantly smaller, as expected, but for the PID control problems, even a few percent improvement means a significant improvement for the system (depending on the selected evaluation function). Details of the tested control systems can be found, e.g. in [25].

The best base results were achieved for the MNIA algorithm, and the proposed mechanisms allowed for an additional improvement of 152%. The greatest improvement of results by the stagnation detection and prevention mechanisms was achieved for the MPGWO algorithm - 217% improvement compared to the base approach - see Table 6. Nevertheless, the use of the MNIA algorithm with different population-based algorithms as islands gives very satisfactory results and at the same time eliminates the need to choose a specific population-based algorithm - thus it is a more universal solution. Stagnation detection improved most of the simulation variants, with the worst improvement being achieved for the MPCS algorithm, with an average improvement of 107% - see Table 6.

Regardless of the approach, algorithm, and strategy, the proposed stagnation detection criterion achieved the best results (see ELA in Table 7). In the case of the best-accuracy strategy (IA), LDFWB worked best (see LDFWB in Table 7), while the complexity-oriented strategy (RC) also achieved an improvement in accuracy - in this case, the LDRWB mechanism turned out to be the best (see LDRWBin Table 7).

The influence of the parameters  $\alpha$ ,  $\beta$ , and  $\gamma$  of the proposed ELA approach and *Nimp* of the SFF approach may be of key importance for various simulation problems and is a topic worth addressing in future research - see Figure 3. Research on other Table 6. Results averaged across all benchmark functions and migration topologies. The results show the improvement in results (in percent) relative to the same methods without the use of stagnation detection and prevention mechanisms. SM stands for stagnation mechanisms, SS stands for stagnation strategy, BASE stands for averaged normalized results, and allows comparing which topology is best (lower values stand for better results). The best results for the accuracy approach are in bold, and for the complexity reduction approach are underlined.



<b>SM/SS</b>	ISBM	<b>LDTC</b>	<b>TTRC</b>	<b>IRBM</b>	<b>RWAC</b>	<b>SMWD</b>	<b>AVG</b>
<b>STA</b>	189	143	132	158	147	143	152
<b>SFF</b>	129	132	77	169	154	130	132
ELA	203	212	138	177	196	185	185
<b>DBDSW</b>	305	191	139	366	202	198	233
<b>DWDSB</b>	196	108	50	169	27	89	106
<b>LWIBS</b>	180	146	139	178	201	203	175
<b>LBIWR</b>	385	377	371	63	109	83	232
<b>LDFBW</b>	119	169	150	267	254	193	192
<b>LDFWB</b>	248	395	8	169	500	376	283
<b>DSBST</b>	37	39	75	49	56	62	53
<b>DSWRS</b>	77	57	63	133	65	67	77
<b>LWRWR</b>	111	99	88	149	125	115	115
<b>LWRBS</b>	129	127	101	166	148	147	136
<b>LDRBW</b>	123	116	90	154	134	123	123
LDRWB	170	123	113	155	168	176	151

Table 7. Summary of tested mechanisms averaged for all benchmark functions. The results show the improvement in results (in percent) relative to the same methods without the use of stagnation detection and prevention mechanisms. SM stands for stagnation mechanisms, SS stands for stagnation strategy. The best results are in bold.

stagnation detection mechanisms, in particular hybrid mechanisms, is also worth considering and is not excluded in the future.

ideologies, which is criticized, and at the same time provide a further field for the development of research in which other similar solutions can be used.

# 5 Conclusions

Mechanisms of stagnation detection and prevention allow for achieving better results in the case of multi-population-based algorithms in most cases. In particular, the best results in averaging the tested topologies, algorithms, simulation problems, and stagnation prevention mechanisms were achieved for the proposed stagnation detection approach (ELA). This approach is based on the analysis of the local improvement of the evaluation function and the infinite impulse response filter. The considered mechanisms for the appropriate management of populations and their sizes also contributed to the improvement of results and additionally enabled the simultaneous reduction of computational complexity. A significant improvement in the results was also achieved for the MNIA algorithm, which uses different baseline population algorithms within the islands. The achieved results point to the possibility of developing multipopulation-based algorithms without the use of new

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