

Doctoral Thesis

Boundary Constraint Violation in Evolutionary Algorithms

Porušování Limitů Argumentů v Evolučních Algoritmech

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"Live slow, live well." – Vincent Van Sloth

ABSTRAKT

V posledních desetiletích se evoluční algoritmy (EA) staly populárními a uznávanými pro svou robustnost a efektivitu v řešení rozmanitých optimalizačních problémů. S narůstajícími výzvami v oblasti umělé inteligence (AI), zejména v kontextu aplikací strojového učení, dochází k nové vlně výzkumu EA. Klíčové aspekty pro další generaci těchto algoritmů zahrnují teoretické základy, analýzy běhu, správné benchmarkovací postupy a detailní zvládání kritických situací, což jsou základní stavební kameny pro dosahování nových úspěchů v AI. Jednou z klíčových výzev je i zvládnutí limitů parametrů optimalizované úlohy, které definují prostor přípustných řešení. I když se dostupné publikace, zabývající se metodami zabraňujících překročení těchto limitů, postupně zlepšují a jejich počet roste, stále je to mnohdy opomíjené téma, které má významný dopad na efektivitu evolučních algoritmů (EA). Tato dizertační práce se zaměřuje na vliv různých protiopatření na výkon evolučních algoritmů (EA). Výzkum začal analýzou základních variant EA, jako jsou PSO (Particle Swarm Optimization), FA (Firefly Algorithm) a SOMA (Self-Organizing Migrating Algorithm). Pozornost se poté přesunula na pokročilejší algoritmy (state-of-the-art), vybrané na základě benchmarkových sad. Studie identifikovala, že integrace účinných protiopatření do návrhu algoritmů může významně ovlivnit jejich pozici v benchmarkových testech. Závěry práce poukazují na významnou problematiku v replikovatelnosti algoritmů, způsobenou nekompletními popisy v publikacích. Tato situace naznačuje potřebu zlepšení v procesu návrhu algoritmů, aby se zvýšila jejich ověřitelnost a udržitelnost.

SUMMARY

In recent decades, evolutionary algorithms (EA) have become popular and recognized for their robustness and effectiveness in solving a variety of optimization problems. With increasing challenges in the field of artificial intelligence (AI), especially in the context of machine learning applications, a new wave of EA research is emerging. Key aspects for the next generation of these algorithms include theoretical foundations, runtime analyses, proper benchmarking procedures, and detailed handling of critical situations, which are fundamental building blocks for achieving new successes in AI. One of the key challenges is mastering the limits of parameters of the optimized task, which define the space of permissible solutions. Although the available publications dealing with methods to prevent exceeding these limits are gradually improving and increasing in number, it is still a neglected topic that has a significant impact on the effectiveness of evolutionary algorithms (EA). This dissertation thesis focuses on the impact of various countermeasures on the performance of evolutionary algorithms (EA). The research began with an analysis of basic variants of EA, such as PSO (Particle Swarm Optimization), FA (Firefly Algorithm), and SOMA (Self-Organizing Migrating Algorithm). Attention then shifted to more advanced algorithms (state-of-the-art), selected based on benchmark sets. The study identified that integrating effective countermeasures into the design of algorithms could significantly influence their position in benchmark tests. The conclusions of the work point to a significant issue in the replicability of algorithms, caused by incomplete descriptions in publications. This situation indicates the need for improvement in the algorithm design process to enhance their verifiability and sustainability.

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1 INTRODUCTION

In recent decades, the landscape of optimization problem-solving has been significantly transformed by the advent and proliferation of metaheuristic algorithms, which have become indispensable tools for tackling tasks of varying complexity across both real and discrete domains. These methods, encompassing a diverse range of techniques from deterministic approaches [\[1\]](#page-73-1) like the Newton [\[2\]](#page-73-2) and gradient methods [\[3\]](#page-73-3) to Evolutionary Algorithms (EAs) [\[4,](#page-73-4) [5,](#page-73-5) [6\]](#page-73-6) and beyond, offer a robust framework for addressing challenges that are often intractable through traditional deterministic means due to current computational limitations. EAs, a prominent subgroup within the metaheuristic category [\[7,](#page-73-7) [8,](#page-73-8) [9\]](#page-73-9), stand out for their effectiveness in solving complex optimization problems, thereby highlighting the vast potential of Evolutionary Computing Techniques (ECTs).

The evolution of metaheuristic algorithms involves expanding theoretical foundations, enhancing strategies through hybridization [\[10\]](#page-73-10), hyperparameter finetuning [\[11\]](#page-73-11), and the integration of learning and adaptive mechanisms. Recent developments also focus on the automatic configuration and selection of algorithms [\[12\]](#page-74-0), merging machine learning with evolutionary computation for improved performance prediction [\[13\]](#page-74-1) and explainability [\[14\]](#page-74-2), highlighting the growing importance of advanced ML techniques in algorithm optimization.

The significance of these advancements is often validated through rigorous benchmarking [\[15\]](#page-74-3) on well-recognized platforms like the IEEE CEC benchmarks [\[16\]](#page-74-4) and the COCO platform [\[17\]](#page-74-5) BBOB testbed, which play a pivotal role in shaping the direction of algorithm design, comparison, and refinement.

Amidst this progress, the metaheuristic research community has increasingly focused on establishing best practices for benchmarking [\[18\]](#page-74-6) to ensure fair, comprehensive comparisons and to gather deeper insights into algorithmic performance across diverse problem sets [\[19\]](#page-75-0). The push towards more transparent and standardized benchmarking protocols reflects a broader recognition of the need to enhance the reproducibility and clarity of results, particularly as metaheuristic algorithms are tasked with solving ever-more complex optimization problems. This initiative, supported by both independent researchers and professional organizations like IEEE, underscores the critical importance of a thorough understanding of optimization processes and methodologies, not only to advance the current state of knowledge but also to facilitate the transition of these insights into other advanced Artificial Intelligence (AI) methods.

This dissertation thesis delves into the crucial aspect of metaheuristics design and benchmarking: the Boundary Control Methods (BCM). Given the inherent variability and randomness of metaheuristic algorithms, there is always a possibility that trial solutions might fall outside the predefined parameter boundaries. This occurrence presents a significant challenge as these boundaries reflect the constraints of real-world optimization scenarios or follow the specification of benchmark functions. Such boundaries might be necessitated by practical considerations, such as ensuring the length of a screw remains within positive numerical limits, or by the theoretical constructs underlying benchmark functions, which dictate constraints to maintain mathematical validity in optimization research.

Boundary limits, therefore, form a core component of virtually every optimization task, necessitating the deployment of effective BCMs to handle instances where trial solutions fall outside these acceptable ranges. Over the years, a diverse array of strategies has been developed to address boundary violations, with some methods gaining prominence for their general applicability across a wide spectrum of algorithms, while others are finely tuned to the nuances of specific problems or algorithms.

By focusing on the intricacies of BCMs, this thesis endeavors to offer valuable insights and recommendations to researchers in the field of metaheuristics, potentially informing future directions in the profiling of benchmark testbeds. The ability to effectively manage boundary constraints is pivotal, not only for the integrity of the optimization process but also for the broader applicability and relevance of metaheuristic algorithms in tackling complex optimization tasks, whether they are rooted in theoretical challenges or practical applications. This

exploration underscores the importance of BCMs as a critical factor in the design, evaluation, and enhancement of metaheuristic algorithms.

The thesis aims to highlight the significance of BCMs as crucial hyperparameters within metaheuristics. These methods should not only undergo thorough optimization and selection but also be clearly documented within the algorithm's description. Such transparency is essential for ensuring the reproducibility of benchmarking outcomes and enhancing our comprehension of metaheuristic population dynamics.

The subsequent sections will provide concise overviews of various basic metaheuristic algorithms selected for examination and comparison regarding their response to BCMs. These algorithms have been specifically chosen to cover a broad spectrum of strategies, some of which draw inspiration from natural phenomena, swarm behaviors, or self-organization processes [\[20,](#page-75-1) [21,](#page-75-2) [22\]](#page-75-3). This focus on a diverse set of metaheuristics is intentional, addressing a gap in existing research, which typically concentrates on a single algorithm or a closely related family of algorithms. Hence, this effort is directed towards offering a more holistic analysis of boundary constraint methods across the field of evolutionary algorithms, aiming to contribute to a comprehensive understanding of how different BCMs impact a wide range of metaheuristic approaches.

2 GOALS OF THE DISSERTATION THESIS

During my scientific activities, I discovered that only limited attention is given to the use of Boundary Control Methods (BCM). Therefore, I have set the following goal for this dissertation:

To document and experimentally verify the impact of using various BCMs on the performance of state-of-the-art evolutionary algorithms. Based on the analysis of the test results, I formulated recommendations for good practices in benchmarking of evolutionary algorithms.

Therefore, the essential steps to goal fulfillment were as follows:

- 1. Survey the current state of Boundary Control Methods (BCMs) used in evolutionary algorithms.
- 2. Investigate influence of various BCMs on the performance of selected evolutionary algorithms.
- 3. Conduct experiments evaluating the impact of BCMs on the performance of state-of-the-art algorithms and results of competitive benchmarking.
- 4. Based on the experimental results draw conclusions and recommendations for good practices in benchmarking.

3 EVOLUTIONARY ALGORITHMS

The motivation behind Evolutionary Algorithms (EAs) [\[4,](#page-73-4) [5,](#page-73-5) [6\]](#page-73-6) began with a relatively simple idea: nature, over millions of years, has evolved systems and organisms that are remarkably effective at solving complex problems. Examples include ants that intuitively find the shortest paths [\[23\]](#page-75-4), bees optimizing their foraging strategies [\[24\]](#page-75-5), the optimal growth patterns of certain plant species [\[25\]](#page-75-6), and various species adapting to environmental changes through natural selection. The foundation of EAs is built upon Charles Darwin's theory of natural evolution [\[26\]](#page-75-7), Gregor Mendel's research on species heredity [\[27\]](#page-75-8), and studies on collective behavior and patterns [\[28\]](#page-75-9).

Thus, a straightforward approach is to apply the vast knowledge of natural systems to real-life optimization challenges. In the context of EAs, this means transforming these natural experiences into metaheuristic algorithms. The general process often involves taking inspiration from a natural process, which is then translated into the pseudo-code of a new algorithm. However, this translation often involves simplification and metaphorization, leading to algorithms that may not always be as robust as their natural counterparts. Moreover, despite the wide array of algorithms within the EA field, only a few have achieved widespread popularity and use. The rapid proliferation of metaheuristic algorithms has drawn criticism [\[29,](#page-76-0) [30\]](#page-76-1), particularly regarding the development of new algorithms based on unjustified concepts drawn from nature, created primarily for rapid publication.

Growing consensus indicates that the creation of these algorithms, despite their innovative surface, often lacks substantial justification in terms of the natural concepts they are based on. Many in the field express concern that such algorithms are more about achieving publication than meaningfully contributing to the field of optimization. This practice not only dilutes the impact of genuinely innovative algorithms but also raises questions about the effectiveness and robustness of these rapidly produced methods. Therefore, there is a pressing need for more rigorous standards in the development and evaluation of new EAs.

These standards should prioritize not just the novelty of the natural concept but also the practical applicability and improvement in optimization effectiveness, ensuring that contributions to the field are both scientifically sound and technically valuable, aligning more closely with the principles of meaningful scientific advancement rather than the pursuit of publication metrics.

3.1 Principles and Mechanisms of Evolutionary Algorithms

In the realm of computational problem-solving, optimization plays a pivotal role, particularly within the context of Evolutionary Algorithms (EAs) [\[31\]](#page-76-2). Optimization fundamentally involves selecting the most effective combination of parameters. This combination is then evaluated based on its overall performance, typically measured by an objective function. The mathematical goal is to find a solution $\mathbf{x} = \{x_1, x_2, \dots, x_D\}$ that optimizes (minimizes or maximizes) the objective function $f(x)$. This optimization process can be subject to various constraints, either defined by functions $g_k(x) \leq 0$; $k = 1, \ldots, n$ or by parameter bounds $l_j \leq x_j \leq h_j$; $j = 1, ..., D$, where D represents the dimensionality of the optimization problem, indicating the number of parameters to be optimized.

EAs approach this optimization task by simulating the process of natural evolution. They operate on a population of potential solutions, applying mechanisms akin to genetic inheritance, mutation, recombination, and selection. Through iterative processes, these algorithms evolve the population, aiming to improve the quality of solutions with respect to the objective function. Each member of the population, often referred to as an individual or a chromosome, represents a potential solution to the optimization problem, encoded in a structure analogous to the genetic makeup of living organisms.

The effectiveness of EAs in navigating complex, multidimensional search spaces lies in their ability to balance exploration and exploitation [\[32\]](#page-76-3). Exploration involves searching through new, untested areas of the solution space, while exploitation focuses on refining the already discovered promising areas. This balance is critical in avoiding local optima traps and steering the algorithm towards the global optimum.

In summary, the essence of optimization in the context of EAs is rooted in the careful selection and iterative improvement of solutions, guided by the principles of natural evolution. This approach is particularly effective in dealing with complex problems where traditional optimization techniques may struggle. Traditional methods often face challenges with complex problems due to their tendency to become trapped in local optima, their sensitivity to the dimensionality of the search space, and their inability to handle discontinuous or nondifferentiable functions effectively. Unlike these methods, EAs are more robust in exploring diverse solution spaces, as they use populations of solutions and stochastic processes, which help avoid premature convergence on sub-optimal solutions and provide a broader exploration of the problem space.

3.2 Genetic Algorithm

The Genetic Algorithm (GA), a renowned member of the EAs family, was pioneered in the 1960s by John Holland [\[33\]](#page-76-4). A hallmark of the classical GA is its unique approach to representing individual parameters encoded as a binary string. These parameters, termed genes, collectively form a chromosome, representing one individual's solution. An entire set of these chromosomes constitutes the population in GA. The main functional components of GA are segmented into three distinct phases.

Selection: This initial phase involves choosing parent chromosomes from the population for reproduction, based on their performance as measured by the objective function. Common methods employed for selection include the roulette rule, rank selection, and tournament selection. Each method influences the rate of convergence and the overall quality of the resulting solution. Typically, individuals with higher solution quality have a greater probability of being selected to pass their genetic information to the subsequent generation.

Crossover: This phase is central to the genetic algorithm, epitomizing the core concept of genetic recombination. The crossover process entails combining genetic information from two parent chromosomes to produce offspring. Various crossover methods exist, but the focus here is on the single-point crossover. As the name implies, single-point crossover involves dividing each parent's chromosome at a specific point and exchanging the segments, resulting in two offspring that inherit a blend of genetic material from both parents.

Mutation: The final phase is mutation, where, with a low probability, certain genes in the offspring's chromosomes may be inverted. The primary objective of mutation is to introduce genetic diversity into the population and to avert premature convergence on suboptimal solutions. The offspring, post-mutation, are then included in the new generation for the next cycle of the algorithm.

Upon completion of the mutation phase, the selection process recommences, marking the start of a new iteration. This cycle continues until a new generation of the population is formed. The description here encapsulates one of the most basic variants of GA. Over the years, numerous adaptations of GA have been developed, many tailored to address specific optimization problems with enhanced efficacy.

The pseudo-code of the described basic GA is shown in Algorithm [1.](#page-19-0)

3.3 Particle Swarm Optimization

Particle Swarm Optimization (PSO) stands as a notable example of so-called swarm-based algorithms, first introduced by Eberhart and Kennedy in 1995 [\[34\]](#page-76-5). The PSO algorithm is inspired by the social behavior and movement patterns observed in natural swarms, such as bird flocking. According to the authors of the proposed modifications, modern adaptations of PSO primarily focus on addressing its known weakness: the tendency for premature convergence to local optima [\[35,](#page-76-6) [36\]](#page-76-7).

In PSO, each member of the swarm, denoted as x , navigates the solution space of the optimization problem. This movement is governed by two key factors: the individual's current position x and its velocity v. The position of each swarm member is updated according to the equation:

$$
x_{i,j}^{k+1} = x_{i,j}^k + v_{i,j}^k
$$
\n(3.1)

where $x_{i,j}^{k+1}$ represents the position of the *i*-th individual in the *j*-th dimension during the $(k + 1)$ -th iteration of the algorithm.

Each individual in the swarm maintains a record of its personal best position, pBest, while the collective swarm retains information about the overall best solution found, **gBest**. These memories, **pBest** and **gBest**, are integral to the calculation of the velocity v , as described in the following equation:

$$
v_{i,j}^{k+1} = w \cdot v_{i,j}^k + c_1 \cdot r_1 \cdot (pBest_{i,j} - x_{i,j}^k) + c_2 \cdot r_2 \cdot (gBest_j - x_{i,j}^k)
$$
(3.2)

Here, w denotes the inertia weight, c_1 and c_2 are cognitive and social factors, and r_1 and r_2 are random numbers drawn from an unimodal distribution within the range $< 0, 1 >$. The pseudo-code for PSO is detailed in Algorithm [2.](#page-21-1)

3.4 Differential Evolution

Differential Evolution (DE), a notable algorithm introduced by Storn and Price in [\[37\]](#page-76-8), forms the foundation of a family of algorithms widely recognized for their effectiveness in continuous optimization. The majority of DE variants adhere to the foundational principles outlined in the original 1995 publication [\[38\]](#page-76-9).

At its core, DE evolves from a randomly generated population of individuals, where each individual x signifies a potential solution to the optimization problem. These individuals are represented as vectors of parameters, with each component of the vector corresponding to a distinct attribute of the problem. The evolution of these individuals is facilitated through a combination of mutation, crossover, and selection processes, aiming to produce superior offspring in subsequent iterations. This evolutionary cycle continues until a predefined stopping criterion is met, ensuring that each new iteration comprises solutions that are either better or equivalent to those in the preceding iteration. The DE algorithm is fundamentally structured into three key steps.

Mutation: This step involves the formation of a mutated vector \boldsymbol{v} through the combination of vectors from selected individuals. This process is governed by a scaling factor F , as illustrated in the equation:

$$
v_{i,j}^k = x_{r_1,j}^k + F \cdot (x_{r_2,j}^k - x_{r_3,j}^k) \tag{3.3}
$$

Here, $v_{i,j}^k$ represents the mutated vector in the j-th dimension for the *i*-th individual at iteration k. The indices r_1, r_2, r_3 are randomly selected from the population and are distinct from each other and also distinct from i-th individual, with NP denoting the total population size.

Crossover: The creation of the trial vector \boldsymbol{u} occurs in this step. It involves choosing attributes either from the mutated vector \boldsymbol{v} or the original vector \boldsymbol{x} , based on a crossover probability determined by the crossover rate CR. The process is encapsulated in the following equation:

$$
u_{i,j}^{k+1} = \begin{cases} v_{i,j}^k, & \text{if } (rand_j \leq CR) \\ x_{i,j}^k, & \text{otherwise} \end{cases}
$$
 (3.4)

The trial vector \boldsymbol{u} is constructed ensuring that it incorporates at least one parameter from the mutated vector v .

Selection: In this final step, the algorithm selects the superior solution between the trial vector \boldsymbol{u} and the original vector \boldsymbol{x} , based on their respective objective function values $f(x)$ and $f(u)$. The selected solution then proceeds to the next iteration.

Through these systematic steps, DE efficiently navigates the solution space, ensuring the continuous improvement of solutions towards optimal results.

The pseudo-code for the general DE is shown in Algorithm [3.](#page-23-0)

3.5 Self-organising Migrating Algorithm

The Self-Organizing Migrating Algorithm (SOMA), developed by Zelinka in 1999 [\[39,](#page-77-0) [40\]](#page-77-1), represents a significant contribution to swarm-based algorithms, incorporating elements of self-organization and various innovative techniques. Influenced by other EAs, SOMA integrates techniques such as discrete perturbation, akin to the mutation process found in evolution strategies. This aspect of SOMA

not only enhances its effectiveness but also facilitates scalability and adaptability across diverse search spaces.

Central to SOMA is the concept of cooperation among individuals, mirroring the collaborative dynamics observed in PSO. In this context, an individual x represents a potential solution to the optimization problem at hand. The essence of this cooperation, as defined by the algorithm's creator, is the migration process of an individual towards another member within the population, expressed mathematically as:

$$
x_{i,j}^{k+1} = x_{i,j}^k + (x_{L,j}^k - x_{i,j}^k) \cdot t \cdot PRTVector_j \tag{3.5}
$$

In this equation, $x_{i,j}^{k+1}$ denotes the new position of the *i*-th individual in the j-th dimension for the next iteration step $k+1$, with $x_{i,j}^k$ being its current position. The term $x_{L,j}^k$ refers to the position of a 'leader,' selected based on a specific SOMA strategy, and t parameterizes individual discrete steps towards this leader, influenced by user-defined parameters $PathLength$ and $Step$.

The mutation process in SOMA is emulated through the $PRTVector_j$, which is generated for each step t and dictates the dimensions in which an individual will migrate towards the leader. The probability of mutation in each dimension is controlled by a user-defined parameter prt , influencing the mutation strength during migration, as detailed in the following equation:

$$
PRTVector_j = \begin{cases} 1, & \text{if } (rand_j < prt) \\ 0, & \text{otherwise} \end{cases}
$$
 (3.6)

SOMA is characterized by its various strategies for leader selection, with the two primary ones being the 'All-To-One' and 'All-To-All' strategies. The 'All-To-One' strategy (the pseudo-code is shown in Algorithm [4\)](#page-24-1) involves selecting one leader per migration cycle based on their objective function value, with other individuals migrating towards this leader. The 'All-To-All' strategy, conversely, sees each individual migrate towards all others in succession, returning to its original position after each migration. The effectiveness of SOMA is showcased in these diverse strategies, each contributing uniquely to the algorithm's ability to explore and exploit the solution space effectively.

Algorithm 4 SOMA - All-To-One

```
1: SOMA initialization
2: while Stopping criterion not met do
3: select leader x_L from population
4: for i = 1 to NP do
5: for t = Step \text{ to } pathLength \text{ with } t+=Step \text{ do}6: generate PRTVector(3.6)
7: migrate x_i to x_L(3.5)
8: evaluate the migrated x_i9: end for
10: save best x_i to the new population
11: end for
12: record the best solution
13: end while
```
3.6 Current Trends in Evolutionary Algorithms

The widespread use of Evolutionary Algorithms, particularly swarm-based ones, presents a paradox: their popularity has been both beneficial and problematic, prompting criticisms for the swift proliferation of new algorithms [\[29\]](#page-76-0). Despite these criticisms, research indicates [\[41\]](#page-77-2) that the development of new algorithms continues steadily, affirming that the trend of creating innovative algorithms is ongoing. Consequently, there is a growing focus among experts on exploring deeper aspects of metaheuristic algorithms, such as runtime analysis, benchmarking, communication dynamics, diversity management, population stagnation, and robustness and efficiency studies [\[42,](#page-77-3) [43,](#page-77-4) [44,](#page-77-5) [45,](#page-77-6) [46,](#page-77-7) [47,](#page-77-8) [48,](#page-77-9) [49,](#page-78-0) [50\]](#page-78-1). This shift is crucial because, despite their practical efficacy, metaheuristic algorithms require a stronger theoretical foundation to fully leverage their capabilities and facilitate the development of new technologies.

This dissertation is aligned with this emerging focus and aims to deliver a detailed examination of a frequently neglected aspect of metaheuristic algorithms—the boundary control methods (BCM). The reproducibility of research is fundamental; thus, omitting critical details renders such studies impractical. Given that BCM can significantly influence the performance and other behavioral aspects of algorithms, it is vital to address them with the same rigor as all other control variables within these algorithms.

4 BOUNDARY CONTROL METHODS

The spectrum of optimization problems is vast, often originating from a wide array of real-world challenges. Typically, these problems are transferred to mathematical forms to facilitate easier analysis. With growing interest in metaheuristic optimization [\[29\]](#page-76-0), there has been a notable increase in the number of benchmark functions and artificial problems created for testing purposes. Each of these benchmark functions used to evaluate metaheuristic optimizers is defined within a specific domain, such as real numbers, positive numbers, or integers, reflecting the varied nature of optimization scenarios. A common feature across both real-world and artificial optimization tasks is the presence of parameter bounds. These bounds may originate from practical limitations in real-world scenarios, such as physical constraints, cost factors, or time limitations.

Due to the inherent randomness in metaheuristic algorithms, there is always a possibility that trial solutions might fall outside the predefined parameter boundaries. This occurrence poses a challenge in effectively solving optimization problems. A typical solution involves checking each newly generated solution to ensure it remains within the acceptable parameter bounds. If a solution is found outside these bounds, an appropriate correction mechanism must be employed to bring it back into the feasible solution space.

The literature presents various methods for addressing boundary constraint violations. Studies suggest that the choice of BCM might influence the performance and other characteristics of an optimization algorithm, as suggested by general studies mentioning potential impacts, often focused on single algorithms. Despite this, there appears to be a lack of comprehensive, generalized studies examining the impact of BCMs on the performance of metaheuristic optimizers in existing publications.

4.1 Literature Overview

The exploration of boundary control methods is addressed in various research publications, with a range of depth in problem coverage. The body of literature pertaining to EA, specifically within the realm of metaheuristic optimization, can be categorized as follows:

- papers primarily dedicated to the exploration of BCMs as a fundamental aspect of EA,
- articles that acknowledge the use of such methods (often within the framework of novel algorithm design) or that develop/implement these methods in a specific optimization algorithm,
- and studies that neglect to discuss BCMs, though inclusion might be beneficial.

It is important to note that searching for relevant literature on this topic is complicated by the similarity in terminology between parameter bounds and objective function constraints. Frequently, while objective function constraints are the focus, parameter bounds are also examined. Thus, thoroughly investigating all related research, which encompasses numerous detailed studies, is both time-intensive and complex.

The primary category of papers should be those entirely dedicated to boundary control handling as a general topic within EA, encompassing various families of metaheuristic algorithms. Regrettably, there are few, if any, comprehensive resources in the literature that address the full spectrum of metaheuristic algorithms.

Papers that discuss boundary control techniques exhibit a range of focuses. While some papers concentrate solely on comparing these methods within a single algorithm, accompanied by statistical analyses and recommendations on which methods generally enhance algorithm performance, the conclusions of these studies are often specific to the algorithm under consideration. The proposed BCMs typically exploit unique aspects of these algorithms, which may limit their applicability for general use.

For PSO-based algorithms, the experimental analysis by Helwig, Branke, and Mostaghim in 2012 [\[51\]](#page-78-2) compares several BCMs within the PSO algorithm. They concluded that such methods significantly affect algorithm performance and can introduce a considerable search bias. This analysis also considered various aspects of PSO, such as the velocity of individual particles and variables for optimal positions. Oldewage, Engelbrecht, and Cleghorn conducted a similarly comprehensive study on BCMs for the PSO algorithm in 2018 [\[52\]](#page-78-3), finding that the hyperbolic method performed best, though its application was limited exclusively to PSO. Clerc's slightly less detailed research in 2006 [\[53\]](#page-78-4) categorized BCMs in PSO as "confinements," describing and testing several on a limited dataset. Zhang et al. conducted a parallel study in 2004, comparing a different set of methods [\[54\]](#page-78-5). Additionally, Michalewicz and Koziel published an extensive study on parameter bounds mixed with constrained numerical optimization for GAs [\[55,](#page-78-6) [56\]](#page-78-7). Detailed work on infeasible solutions and five BCM strategies for addressing them in terms of DE are discussed by Kononova et al. [\[57\]](#page-78-8).

Another subset of articles that mention BCMs do so only as part of the design of new metaheuristic algorithms, with minimal to no analysis. For instance, in their seminal effort to define a standard for PSO, Bratton and Kennedy [\[58\]](#page-79-0) in 2007 only briefly mention the boundary conditions, selecting one type without any detailed discussion or references. Similarly, the paper by Mostaghim et al. in 2006 [\[59\]](#page-79-1) focuses on the multi-objective version of PSO. While primarily dedicated to objective function constraints, it also summarily addresses BCMs alongside previous work. However, extracting relevant results from this study is somewhat challenging due to its primary focus. In another context, four BCMs are discussed in Hansen's tutorial paper on the Covariance Matrix Adaptation Evolution Strategy (CMA-ES) algorithm [\[60\]](#page-79-2), yet the impact of these methods on CMA-ES performance was not explored in the tutorial.

Up to this point, the majority of referenced research articles have focused on the family of PSO algorithms, with one notable exception related to CMA-ES.

This category of papers also encompasses research in the field of Differential Evolution (DE) algorithms. Ronkkonen et al. [\[61\]](#page-79-3) describe a method for the DE algorithm where a trial solution is reflected off the boundary by the amount of the violation. A similar technique is employed in the works of Brest et al. [\[62\]](#page-79-4), Price and Storn [\[63\]](#page-79-5), Guo et al. [\[64\]](#page-79-6), and Zhang et al. [\[65\]](#page-79-7). Additionally, Caraffini et al. provide more detailed work on structural bias—primarily caused by boundary constraints—with extensive results and discussion in [\[66\]](#page-79-8). Further work by Vermetten et al. led to the creation of a toolbox called BIAS, designed specifically for benchmarking structural bias, as detailed in [\[67\]](#page-80-0).

The results and implications derived from the collected works suggest that BCMs can directly influence the overall performance of a metaheuristic algorithm. Yet, many new algorithms, tutorials, and overview articles have overlooked or neglected this aspect. For instance, this oversight occurs in a tutorial for PSO [\[68\]](#page-80-1), a general paper on EAs [\[69\]](#page-80-2), articles on the Firefly Algorithm (FA) [\[70,](#page-80-3) [71\]](#page-80-4), a publication on the Cuckoo Algorithm [\[72\]](#page-80-5), and a paper introducing new mechanics for DE [\[73\]](#page-80-6).

The following quotation from the article on experimental analysis of BCM (stated as bound handling techniques in the article) in particle swarm optimization by Helwig et al. [\[51\]](#page-78-2) emphasizes this point:

"As was shown, the bound handling technique has a significant impact on the performance of PSO, particularly when the search space is of high dimensionality, as this greatly increases the likelihood of a particle exiting the feasible area."

This statement, along with other publications listed in subsection [4.1,](#page-27-0) strongly suggests that BCMs require careful consideration—whether in proposing a new metaheuristic algorithm or when modifying an existing one. The findings in these publications also indicate that different methods are preferred for different algorithms, underscoring the need for extensive research to examine the employed methods across a variety of optimizers. Such research should aim to deeply analyze their effects not only on performance but also on other characteristics of the algorithms.

4.2 Existing Boundary Control Methods

This subsection presents a consolidated overview of Boundary Control Methods (BCMs) commonly utilized in research. The versatility of some techniques allows for their adaptation in various forms, while others are specifically tailored for particular optimization algorithms. Additionally, certain methods might be underrepresented in this summary due to their lesser popularity or because they are briefly mentioned in research focusing on different topics.

In the ensuing subsections, mathematical formulations of various BCMs are detailed. To facilitate comprehension, a brief recapitulation of the variables employed in these formulations is provided here. An individual solution in the context of these methods is denoted as $x = \{x_1, x_2, \ldots, x_D\}$, representing a vector of parameters where D indicates the dimension size. Each parameter x_i within this vector, corresponding to a specific dimension, is subject to defined bounds that demarcate the feasible solution space. These parameter bounds are expressed as $l_j \leq x_j \leq h_j$, where l_j and h_j represent the lower and upper bounds, respectively, for the j-th dimension.

4.2.1 Clipping

The Clipping Method (also known as saturation) stands out for its simplicity and ease of implementation, often making it one of the first choices in BCMs. In this method, individual solutions x are prevented from crossing the defined boundaries in each dimension. Instead, they are "clipped" to remain within the parameter bounds. This approach is succinctly captured by the following equation, ensuring that each solution stays within its maximum and minimum limits.

$$
x_{i,j}^{k+1} = \begin{cases} h_j, & \text{if } \left(x_{i,j}^k > h_j\right) \\ l_j, & \text{if } \left(x_{i,j}^k < l_j\right) \\ x_{i,j}^k, & \text{otherwise} \end{cases} \tag{4.1}
$$

The $x_{i,j}^{k+1}$ is the *i*-th individual in *j*-th dimension in $k+1$ iteration, and the pair h_j and l_j are the parameter bounds, maximum and minimum respectively.

4.2.2 Random

In cases where a trial solution violates the boundary in any dimension, the Random Method generates a new position for the respective dimension. This position is randomly determined between the lower and upper bounds, following a pseudo-random uniform distribution. This method is straightforward to implement, as reflected in its mathematical representation.

$$
x_{i,j}^{k+1} = \begin{cases} U(l_j, h_j) & \text{if } \left(x_{i,j}^k > h_j \text{ OR } x_{i,j}^k < l_j\right) \\ x_{i,j}^k & \text{, otherwise} \end{cases} \tag{4.2}
$$

4.2.3 Reflection

The Reflection Method mirrors a solution back into the feasible space if it attempts to cross the defined borders. This technique is akin to the reflection behavior of a mirror. For each dimension that violates the boundary, the position of the individual is corrected in a way that reflects it back into the permissible range.

$$
x_{i,j}^{k+1} = \begin{cases} h_j - \left(x_{i,j}^k - h_j\right) & \text{if } \left(x_{i,j}^k > h_j\right) \\ l_j + \left(l_j - x_{i,j}^k\right) & \text{if } \left(x_{i,j}^k < l_j\right) \\ x_{i,j}^k & \text{,otherwise} \end{cases} \tag{4.3}
$$

4.2.4 Periodic

The Periodic Method approaches boundary violations by considering an infinite solution space, effectively creating infinite copies of the optimized hyper-space. It employs a mapping technique that brings the individual back into the feasible space using a modulo function. This method ensures that solutions are cyclically repositioned within the acceptable range.

$$
x_{i,j}^{k+1} = l_j + \left((x_{i,j}^k - h_j) \text{ MOD } (h_j - l_j) \right) \tag{4.4}
$$

4.2.5 Halving the Distance

As suggested by its name, this method involves halving the distance between the original position and the crossed boundary. Unlike previous techniques, this approach requires tracking the starting position of an individual. It offers a more nuanced adjustment by averaging the boundary and the initial position.

$$
x_{i,j}^{k+1} = \begin{cases} x_{i,j}^{k-1} + \left(h_j - x_{i,j}^{k-1} \right) / 2 & \text{, if } \left(x_{i,j}^k > h_j \right) \\ l_j + \left(x_{i,j}^{k-1} - l_j \right) / 2 & \text{, if } \left(x_{i,j}^k < l_j \right) \\ x_{i,j}^k & \text{, otherwise} \end{cases}
$$
(4.5)

4.2.6 Soft

The Soft Method is unique in that it imposes no immediate restrictions on individuals outside the feasible space, except that their objective function values are not updated until they re-enter the feasible area. Implementing this method can be challenging, as it does not guarantee finite iteration completion without specific algorithm tuning. This approach allows for greater flexibility but requires careful management to ensure algorithm convergence.

5 CHRONOLOGICAL CONTRIBUTIONS TO BOUND-ARY CONTROL METHODS

The subsequent sections outline the empirical findings from a set of focused studies undertaken by the author of this doctoral thesis. Initial observations concentrated on the Particle Swarm Optimization (PSO) and its advanced iterations; further studies were conducted on the Self-Organizing Migrating Algorithm (SOMA) and the Firefly Algorithm (FA) [\[74\]](#page-80-7). These studies aimed to explore the effect of the Boundary Constraint Methods (BCMs) on the overall performance of these metaheuristic algorithms.

An additional cornerstone of this exploration is presented in the journal article titled "Impact of Boundary Control Methods on Bound-Constrained Optimization Benchmarking" [\[75\]](#page-80-8). This paper focuses on the critical implementation of BCMs and their significant influence on the performance of leading algorithms, as demonstrated in the IEEE CEC competitions of 2017 and 2020. The motivation for this research was based on the findings described in previous initial studies [\[76,](#page-80-9) [77,](#page-81-0) [78\]](#page-81-1), where the influence of BCMs on the performance of selected basic versions of metaheuristic algorithms was examined. Thus, a research question arose as to whether the choice of a BCM could significantly influence other competitive algorithms, especially the CEC competition winners. Furthermore, it was motivating to find out whether just changing the BCM can help achieve even better results for the top three performing algorithms from a given year of the competition, possibly changing their final order. The results presented in the article showed that the impact on the empirical performance of the top 3 algorithms from CEC17 is affected by the choice of BCMs.

A further contribution was focused on exploring the frequency of BCM activations. The study investigated how often BCMs were activated during the optimization process across various metaheuristic algorithms.

A timeline of the contributions mentioned, detailing the progression and key milestones of the research, is illustrated in Figure [5.1.](#page-35-1)

Fig. 5.1 Research timeline progress.

5.1 PSO

The study's results published in 2017 [\[76\]](#page-80-9) confirmed that the selection of used BCM affects the algorithm's performance. The study compared Clipping, Random, Periodic, and Soft methods on the generic version of PSO and the variant called Diversity guided PSO (ARPSO) [\[35\]](#page-76-6). The ARPSO algorithm, developed by J. Riget and J. S. Vesterstrøm in 2002, specifically addresses the issue of premature convergence – a notable shortcoming of the traditional PSO algorithm they highlighted in their proposal.

Operating on principles akin to those outlined in Section [3.3](#page-20-0) for PSO, ARPSO involves computing the population's diversity in each iteration. Should this divergence fall below a predefined threshold, a repulsive phase is initiated to disperse the particles, thereby enhancing exploration. Conversely, if the divergence exceeds a certain upper threshold, an attractive phase is triggered, encouraging
particles to converge, which aids in exploitation. This dynamic adjustment between attraction and repulsion phases helps in maintaining a balance between exploration and exploitation, which is crucial for avoiding premature convergence.

5.1.1 Experimental setup

The experiments were conducted on the CEC 2015 benchmark set [\[79\]](#page-81-0) for dimension sizes 10, 30, and 50. The benchmark encompasses 15 test functions, and every test function was repeated for 51 independent runs. The results were tested for statistical significance using the Friedman Rank test [\[80\]](#page-81-1) with the significance level $\alpha = 0.05$, accompanied by Nemenyi critical distance (CD) [\[81\]](#page-81-2).

5.1.2 Key findings

The Friedman rank tests (Fig. [5.2,](#page-37-0) Fig. [5.3\)](#page-38-0) clearly show that the Clipping method negatively affected the overall performance of both PSO and ARPSO algorithms on all dimension sizes. The CD shows which BCMs do not alter the performance of the particular algorithm with statistical significance in comparison with the first-ranked method. These are Random, Periodic, and Soft methods for dimension sizes 10 and 30. For dimension size 50, only Random and Soft methods perform similarly.

5.2 FA

The subsequent study from 2018 [\[77\]](#page-81-3) compared Clipping, Random, Reflection, and Periodic BCMs on FireFly Algorithm (FA) and on a hybrid of FA and PSO, called Firefly Particle Swarm Optimization (FFPSO) [\[82\]](#page-81-4). This hybrid algorithm was introduced in late 2016 by Padmavathi Kora and K. Sri Rama Krishna. The basic idea behind such an approach, according to the authors, is

Fig. 5.2 Friedman rank comparison of the Clipping, Random, Periodic, and Soft methods on PSO on benchmark set CEC 2015. [\[76\]](#page-80-0)

that the new hybrid strategy can share advantages from both algorithms and hopefully eliminate their disadvantages. The main principle remains the same as in the standard FA, but the equation for firefly motion is slightly changed according to PSO movement.

5.2.1 Experimental setup

The experiments were performed on the CEC 2017 benchmark set [\[83\]](#page-81-5), encompassing 30 test functions, and every test function was repeated for 51 independent runs. The tested dimension sizes were 10 and 30. The results were tested for statistical significance using the Friedman Rank test [\[80\]](#page-81-1) with the significance level $\alpha = 0.05$, accompanied by Nemenyi critical distance (CD) [\[81\]](#page-81-2).

Fig. 5.3 Friedman rank comparison of the Clipping, Random, Periodic, and Soft methods on ARPSO on benchmark set CEC 2015. [\[76\]](#page-80-0)

5.2.2 Key findings

The figures Fig. [5.4](#page-39-0) and Fig. [5.5](#page-39-1) show the aforementioned ranks for FA and FFPSO algorithms, respectively. The lower the rank, the better the performance of the used BCM. Furthermore, the presented Friedman ranks are again accompanied by Nemenyi critical distance (CD), represented as the dashed line, measuring the critical distance from the best BCM (the lowest mean rank).

According to the ranking shown in both Figures (Fig. [5.4](#page-39-0) and Fig. [5.5\)](#page-39-1), the significant impact of the selected BCM is mostly observed on canonical FA. For the hybrid method FFPSO, the results indicate a minimal or negligible impact of the used method. For dimension sizes 10 and 30, the most favorable BCMs appear to be clipping and reflection methods.

Fig. 5.4 Friedman rank comparison of the Clipping, Random, Reflection, and Periodic methods on FA on benchmark set CEC 2017. [\[77\]](#page-81-3)

Fig. 5.5 Friedman rank comparison of the Clipping, Random, Reflection, and Periodic methods on FFPSO on benchmark set CEC 2017. [\[77\]](#page-81-3)

5.3 SOMA

The 2020 study [\[78\]](#page-81-6) delved into the impact of BCMs on the Self-Organizing Migrating Algorithm (SOMA), specifically its All-To-One and All-To-All variants. SOMA is a metaheuristic optimization technique inspired by the social behavior of individuals within a population moving towards better positions or solutions. It is known for its efficacy in navigating complex optimization landscapes. This investigation was prompted by the notable gap in research concerning the interaction between BCMs and SOMA strategies, a gap that this study aimed to bridge.

The study evaluated four distinct BCMs: Clipping, Random, Reflection, and Periodic. This comparative analysis sheds light on how different BCMs can influence the performance and robustness of SOMA variants, contributing valuable insights to the sparse body of knowledge on BCMs' role in enhancing SOMA's optimization capabilities.

5.3.1 Experimental setup

For the experiment, the CEC 2017 benchmark set [\[83\]](#page-81-5) was chosen, encompassing 30 test functions categorized into unimodal, multimodal, hybrid, and composite groups. The experiment focused on dimension sizes of 10 and 30, adhering to the benchmark's stipulation of a maximum of 10,000 function evaluations per dimension. To ensure robustness, each test function underwent 51 independent trials, with the outcomes subjected to statistical analysis.

Regarding the SOMA parameters, they were configured as follows: the population size NP was 100 individuals; the probability of perturbation prt was set to 0.3; the Step parameter was 0.11, and the Path length was fixed at 3. These settings were in line with the recommendations of the original authors [\[39,](#page-77-0) [40,](#page-77-1) [84\]](#page-82-0).

The analysis utilized the Friedman Rank test [\[80\]](#page-81-1) to assess statistical significance, rejecting the null hypothesis of equal means at a 5% significance level.

5.3.2 Key findings

Figure [5.6](#page-42-0) displays the Friedman ranking outcomes for both the All-To-One and All-To-All SOMA strategies across the dimension sizes of 10 and 30, where lower ranks signify superior performance. To compare multiple BCMs, the Nemenyi Critical Distance post-hoc test was employed, determining a critical distance (CD). The critical distance is visually represented by a dashed line extending from the highest-ranked boundary method in the figure.

For the All-To-One strategy, the Random and Periodic methods emerged as notably effective, with the All-To-All strategy showing similar results and the Reflection method also presenting promise. Conversely, the Clipping method was identified as the least effective across strategies. The analysis distinguished two method groups: one (Clipping, Reflection, and Periodic) maintaining consistent individual migration across the search space without losing previous position information, and the other (Random method) facilitating a more stochastic search, potentially aiding SOMA in avoiding local minima traps.

The findings indicate the Random and Periodic methods as particularly promising, with the Random method introducing beneficial stochastic elements to SOMA. In contrast, the Periodic method preserves individuals' movement directions and patterns, offering a more predictable and natural behavior that aligns with original population dynamics.

The least effective method for both SOMA strategies was found to be the Clipping method, potentially due to its restriction of individual movements to the borders of the feasible space, impacting the algorithm's performance negatively in both tested dimension sizes.

Fig. 5.6 Friedman rank comparison of the Clipping, Random, Reflection, and Periodic methods on SOMA on benchmark set CEC 2017. [\[78\]](#page-81-6)

5.4 Analyzing the Impact of Boundary Control Methods on Algorithmic Performance

The impulse for the performance analysis study comes from earlier investigations [\[76,](#page-80-0) [77,](#page-81-3) [78\]](#page-81-6) that explored the impact of BCMs on basic metaheuristic algorithms, raising questions about their potential influence on more competitive algorithms, particularly CEC competition winners. This study investigates whether modifications to BCMs could enhance the performance of top algorithms from recent competitions, potentially altering their ranking outcomes.

5.4.1 Experimental setup

The study was focused on the three top-ranking participants of two benchmark competitions: CEC17 [\[83\]](#page-81-5), and CEC20 [\[85\]](#page-82-1). The goals for both testbeds were to:

- determine which BCM was used by the three winning algorithms,
- examine if there is a better choice of a BCM for a particular algorithm,
- if the algorithms used a different BCM, could it have changed the final order?

Both benchmarks encompass four groups of test functions: unimodal, multimodal, hybrid, and composition functions. An advantageous feature of the CEC benchmark is that all incorporated test functions are defined with equal and static (same values across all dimensions) search ranges for all parameters. The original implementation also supports a shift of the global optimum and rotation of each function.

The description of each benchmark is summarized in the following subsections alongside the descriptions of the three top-ranking algorithms of each benchmark.

Algorithm	BCM	BCM listed
EBOwithCMAR ^[87]	Halving, Clipping	N ₀
LSHADE-cnEpSin [88]	Halving	N ₀
jSO[89]	Halving	Partially
DES [90]	Penalization	Yes
DYYPO ^[91]		N ₀
IDEbestNsize [92]	Reflection	Partially
LSHADE SPACMA [93]		No
MM OED [94]		No
MOS-CEC2013 [95]		No
MOS-SOCO2011 [95]		No
PPSO [96]	Random bounce	Yes
RB-IPOP-CMA-ES [97]		N ₀
TLBO-FL [98]	Clipping	Yes

Tab. 5.1 CEC17 – Algorithm overview

5.4.2 Top 3 best-performing algorithms for CEC17

The testbed CEC17 published in 2016 [\[83\]](#page-81-5) encompasses 30 test functions for dimension sizes of 10, 30, 50, and 100. The following subsections briefly describe the top three performing algorithms according to the official results [\[86\]](#page-82-7). Table [5.1](#page-44-0) contains a list of all participants, including the used BCM, and if the BCM was mentioned in the accompanying paper.

EBOwithCMAR was initially proposed for the CEC17 benchmark and successfully obtained the first position among 11 competitors. The hybrid algorithm is based on the Effective Butterfly Optimizer (EBO) and Covariance Matrix Adapted Retreat Phase (CMAR), which improves the local search capability of EBO. The paper [\[87\]](#page-82-2) does not specify any used BCM; however, upon the analysis of the algorithm code showed that EBOwithCMAR uses two BCMs, Halving for EBO and Clipping for CMAR.

jSO [\[89\]](#page-82-4) represents an improved variant of the iL-SHADE algorithm [\[99\]](#page-83-7) and ranked in second place. The improvement lies predominantly in the new version

Algorithm	BCM	BCM listed
$\overline{\text{IMODE}}$ [102]	Random, Halving	N ₀
AGSK [103]	Halving	N ₀
$j2020$ [104]	Periodic	Yes
Cssin $[105]$		N _o
MP-EEH [106]	Clamping	Yes
RASP-SHADE [107]	Halving	Yes
DISH-XX [108]	Halving	N ₀
jDE100e [109]		No
OLSHADE [110]	Clipping, Halving	Yes
$mpmL-SHADE$ [111]	Halving	Yes
SOMA-CL [112]	Random	No

Tab. 5.2 CEC20 – Algorithm overview

of the mutation strategy. The jSO uses Halving BCM, which is referred in the paper as a "repeat mechanism" without any detailed description or citation.

LSHADE-cnEpSin is the third algorithm used in this study and represents an extension to the LSHADE-EpSin [\[100\]](#page-84-6), which was ranked as the joint winner in the competition IEEE CEC 2016. The enhancement lies in the ensemble of sinusoidal approaches and covariance matrix learning for the crossover operator. The LSHADE-cnEpSin [\[88\]](#page-82-3) ranked third in the CEC17 competition and uses Halving BCM, which is unfortunately not mentioned in the paper by the authors.

5.4.3 Top 3 best-performing algorithms for CEC20

The CEC20 benchmark [\[85\]](#page-82-1) was introduced in 2019 and includes 10 test functions for dimension sizes of 5, 10, 15, and 20. Again, the following subsections briefly describe the top three performing algorithms according to the official results [\[101\]](#page-84-7), and Table [5.2](#page-45-0) contains a list of all participants, what boundary control method they used, and if the BCM was mentioned in the related paper.

IMODE is a Differential Evolution (DE) based algorithm ranked as the winner

in the CEC20 competition. IMODE [\[102\]](#page-84-0) benefits from multiple differential evolution operators, with more emphasis placed on the best-performing operator. The algorithm employs two BCMs: Clipping and Halving, selected randomly each time it is used. Unfortunately, BCMs are not mentioned in the paper.

AGSK is the second-best performing algorithm in the CEC20 competition. AGSK [\[103\]](#page-84-1) is the enhanced version of the Gaining Sharing Knowledge-based algorithm (GSK) [\[113\]](#page-85-5), which uses adaptive settings to its control parameters. The algorithm utilizes Halving BCM; however, the authors did not specify this.

j2020 [\[104\]](#page-84-2) ranked third place in the competition, and it is based on the two self-adaptive DE algorithms: jDE [\[114\]](#page-85-6) and jDE100 [\[115\]](#page-86-0). The used BCM is Periodic, which is described in the paper.

5.4.4 Statistical Evaluation

Friedman rank test was a first step. Each algorithm was tested and evaluated while using different BCMs. The evaluation was performed by Friedman rank test [\[80\]](#page-81-1), and the results are presented in Table [5.3](#page-50-0) for CEC17 and Table [5.4](#page-51-0) for CEC20. The values in cells are the rankings for each algorithm for a particular dimension size. The columns indicate the tested BCM. If the algorithm used a different BCM than the selected (Clipping, Random, Periodic, Reflection, Halving), the column Default was used. Otherwise, this column states the name of the used BCM of the algorithm. The last column contains p-values of the Friedman rank test. The tested significance levels are 0.1, 0.05, 0.01, and 0.001. Each level corresponds to a certain symbol: *, †, **, and *** respectively. Therefore, a symbol represents the significance level of the result. The last row of each algorithm also contains the mean rank (given in bold) across the dimension sizes for a particular BCM. The last column, CD, stands for Nemenyi Critical Difference – if the difference between a pair of BCMs' ranks is higher than the CD value, they are significantly different. For example, in Table [5.3,](#page-50-0) the Periodic BCM in LSHADE-cnEpSin in 10D is significantly different (better according to rank) from Clipping BCM. But the same Periodic BCM is not significantly different from Random BCM.

Table [5.3](#page-50-0) contains more results with statistically significant differences than Table [5.4.](#page-51-0) The likely reasons are that the CEC20 benchmark contains only 10 test functions and lower dimensionality might cause a lower number of BCM use (see [\[51\]](#page-78-0) – low dimensionality leads to a lower probability of creation of an infeasible trial solution), and the top ranking algorithms in the competition are robust and similar in the performance.

5.4.5 Competition Scoring System

The motivation for this second test, using CEC scoring system, was to determine if a change of BCM may cause a change in the order of the algorithms. The CEC scoring system is in detail provided in the technical reports accompanying the CEC benchmarks [\[83,](#page-81-5) [85\]](#page-82-1).

The final score value sorts the algorithms, and the higher score value, the better is the performance of the algorithm. This Score is a sum of two partial scores named Score 1 and Score 2. Score 2 is based on the weighted rank values, and Score 1 is computed from the normalized error values for CEC20 or mean (not normalized) error values for CEC17. Score 1 is computed using equations (5.1) and [\(5.2\)](#page-48-0).

$$
SE = 0.1 \cdot \sum_{i=1}^{29} e f_{10D} + 0.2 \cdot \sum_{i=1}^{29} e f_{30D}
$$

+0.3 \cdot \sum_{i=1}^{29} e f_{50D} + 0.4 \cdot \sum_{i=1}^{29} e f_{100D} \tag{5.1}

$$
Score1 = \left(1 - \frac{SE - SE_{min}}{SE}\right) \cdot 50\tag{5.2}
$$

Where ef is the mean error value for a certain dimension size and SE_{min} is the minimal sum of errors among all algorithms. The Score 2 is then computed based on equations equations [\(5.3\)](#page-48-1) and [\(5.4\)](#page-48-2).

$$
SR = 0.1 \cdot \sum_{i=1}^{29} rank_{10D} + 0.2 \cdot \sum_{i=1}^{29} rank_{30D}
$$

+0.3 \cdot $\sum_{i=1}^{29} rank_{50D} + 0.4 \cdot \sum_{i=1}^{29} rank_{100D}$ (5.3)

$$
Score2 = \left(1 - \frac{SR - SR_{min}}{SR}\right) \cdot 50\tag{5.4}
$$

The final score is then defined as [\(5.5\)](#page-48-3).

$$
Score = Score1 + Score2 \tag{5.5}
$$

The equations $(5.1) - (5.5)$ $(5.1) - (5.5)$ $(5.1) - (5.5)$ describes the computation of the score for CEC17. The same equations are used to compute the CEC20 score but with different dimension sizes.

5.4.6 Results

This subsection presents the results of both experiments performed for benchmarks CEC17 and CEC20. Each test scenario used a different number of independent runs as defined by the used benchmark. CEC 17 testbed defines 51 independent runs, while CEC20 testbed requires 30 independent runs.

The three approaches were used to analyze and represent the results: Friedman rank test, CEC scoring system, and selection of the best performing BCM variant for the algorithms.

The Tables $5.5 a$ – f) contain the Score and rank of BCMs used for a particular algorithm. The default BCM is in bold. From the given results, not once did the default BCM ranked as the best performing variant; therefore, potentially better results for the algorithm may be achieved using the BCM with the highest Score. The parentheses under the score display percentual contribution of each dimensional setting to the score. The significant disproportion in values of Score 1 for dimension size 50 for CEC17 was caused by the last test function f_{30} , Since this disproportion is observed across all tested algorithms and their BCMs, the obtained results are still comparable, however further investigation is needed to find the cause of such behavior.

Friedman ranks suggest that higher dimension size has a more significant impact on the final score, as can be seen in Table [5.5.](#page-49-0) However, the higher impact of higher dimension sizes is also implicit due to the weighting of dimension parts of the score computation in [\(5.1\)](#page-47-0) and [\(5.3\)](#page-48-1).

Tab. 5.5 Score and rank of BCMs used for a particular algorithm. The rank is based on the final Score, which is a sum of partial scores 1 and 2. The default BCM of the algorithm is in bold. The parentheses under the score display percentual contribution of each dimensional setting to the score (CEC17 – ${10D, 30D, 50D, 100D},$ CEC20 - ${5D, 10D, 15D, 20D}.$

c) CEC17 – LSHADE-cnEpSin

Rank	BCM	Score 1	Score 2	Score
1	Random	$5.00E + 01$	$5.00E + 01$	$1.00E + 02$
		(0.5, 1.2, 91.8, 6.5)	(9.2, 20., 30.4, 40.5)	
$\mathcal{D}_{\mathcal{L}}$	Periodic	$4.92E + 01$	$4.78E + 01$	$9.71E + 01$
		(0.5, 1.2, 91.5, 6.8)	(8., 19.9, 28.3, 43.8)	
3	Reflection	$4.60E + 01$	$4.20E + 01$	$8.79E + 01$
		(0.5, 1.1, 92.5, 5.9)	(10.6, 20.8, 30.2, 38.4)	
$\overline{\mathbf{4}}$	Halving	$4.44E + 01$	$4.21E + 01$	$8.65E + 01$
		(3.7, 1.1, 89.7, 5.5)	(10.9, 19.7, 29.9, 39.5)	
5	Clipping	$4.43E + 01$	$3.66E + 01$	$8.08E + 01$
		(0.4, 1.1, 92.7, 5.8)	(10.8, 19.7, 31., 38.6)	

d) CEC20 – IMODE

Rank	BCM	Score 1	Score 2	Score
$\mathbf{1}$	Reflection	$5.00E + 01$	$5.00E + 01$	$1.00E + 02$
		(1.4, 10.9, 34.5, 53.2)	(10.8, 19.7, 33.9, 35.6)	
$\overline{2}$	Halving	$4.95E + 01$	$4.41E + 01$	$9.36E + 01$
		(2.4, 11.1, 32.4, 54.1)	(8.7, 22.4, 30.7, 38.1)	
3	Random	$4.47E + 01$	$4.79E + 01$	$9.27E + 01$
		(4.4, 10.3, 32.7, 52.6)	(10.5, 22., 26.1, 41.5)	
4	Periodic	$4.35E + 01$	$4.27E + 01$	$8.63E + 01$
		(5.5, 7.9, 39.5, 47.)	(10.2, 20.1, 27.3, 42.4)	
5	Default	$4.37E + 01$	$4.19E + 01$	$8.57E + 01$
		(4.3, 15.9, 29.9, 49.8)	(8.7, 18.1, 31.6, 41.6)	
6	Clipping	$4.14E + 01$	$4.39E + 01$	$8.53E + 01$
		(4.1, 8.9, 27.4, 59.6)	(11.3, 17.9, 30.6, 40.2)	

e) CEC20 – AGSK

Rank	Algorithm	Score 1	Score 2	Score
1	EBOwithCMAR	$5.00E + 01$	$5.00E + 01$	$1.00E + 02$
$\overline{2}$	iSO	$4.97E + 01$	$4.33E + 01$	$9.30E + 01$
3	$LSHADE-enEpSin$	$4.68E + 01$	$4.47E + 01$	$9.15E + 01$
$\overline{4}$	LSHADE SPACMA	$4.64E + 01$	$4.47E + 01$	$9.11E + 01$
5	DES	$4.59E + 01$	$4.12E + 01$	$8.71E + 01$
6	MM OED	$4.60E + 01$	$3.62E + 01$	$8.22E + 01$
7	IDEbestNsize	$2.98E + 01$	$2.63E + 01$	$5.61E + 01$
8	$MOS-CEC2013$	$1.89E + 01$	$1.74E + 01$	$3.63E + 01$
9	RB-IPOP-CMA-ES	$3.79E + 00$	$3.21E + 01$	$3.59E + 01$
10	MOS-SOCO2011	$1.11E + 01$	$1.92E + 01$	$3.03E + 01$
11	PPSO	$3.93E + 00$	$1.73E + 01$	$2.12E + 01$
12	DYYPO	$5.93E - 01$	$1.71E + 01$	$1.77E + 01$
13	TLBO-FL	$2.87E - 02$	$1.64E + 01$	$1.64E + 01$

Tab. 5.6 CEC17 – Score – Default BCM

f) CEC20 – j2020

Rank	BCM	Score 1	Score 2	Score
1	Halving	$5.00E + 01$	$4.27E + 01$	$9.27E + 01$
		(0.5, 0.5, 42.1, 56.8)	(9.6, 24.3, 31.1, 35.)	
\mathfrak{D}	Random	$4.19E + 01$	$5.00E + 01$	$9.19E + 01$
		(0., 5., 35.2, 59.7)	(11.3, 14.2, 27.6, 46.9)	
3	Reflection	$4.35E+01$	$4.34E + 01$	$8.69E + 01$
		(0., 5.6, 43.1, 51.4)	(10.9, 21.1, 28.9, 39.2)	
$\overline{\bf{4}}$	Periodic	$3.62E + 01$	$3.59E + 01$	$7.22E + 01$
		(0., 4.3, 33.1, 62.6)	(8.7, 19.8, 31.1, 40.3)	
5	Clipping	$3.68E + 01$	$3.20E + 01$	$6.88E + 01$
		(0., 6., 14., 80.)	(9.9, 19.8, 30.6, 39.7)	

Selection of the BCM was the third and the last step to implement the best-performing BCM variant for the algorithms and check if the final order of the competition would be different. Table [5.6](#page-54-0) and Table [5.10](#page-56-0) contain the Score and rank if the algorithms used their default BCMs. Unfortunately, the complete results of all competitors are available for the CEC17 benchmark only; therefore, Table [5.10](#page-56-0) encompasses only the three top-ranking algorithms of the CEC20 competition. The best BCM for each algorithm was selected according to the ranks in Tables [5.5](#page-49-0) - [5.5.](#page-49-0)

Rank	Algorithm	Score 1	Score 2	Score
1(1)	EBOwithCMAR	$5.00E + 01$	$5.00E + 01$	$1.00E + 02$
2(2)	iSO	$4.96E + 01$	$4.55E + 01$	$9.51E + 01$
3(4)	LSHADE SPACMA	$4.66E + 01$	$4.83E + 01$	$9.49E + 01$
4(3)	$LSHADE-enEpSin$	$4.78E + 01$	$4.25E + 01$	$9.03E + 01$
5(5)	DES	$4.61E + 01$	$4.23E + 01$	$8.84E + 01$
6(6)	MM OED	$4.62E + 01$	$3.74E + 01$	$8.35E + 01$
7(7)	IDEbestNsize	$3.00E + 01$	$2.69E + 01$	$5.69E + 01$
8(9)	RB-IPOP-CMA-ES	$3.81E + 00$	$3.32E + 01$	$3.70E + 01$
9(8)	$MOS-CEC2013$	$1.90E + 01$	$1.78E + 01$	$3.68E + 01$
10(10)	MOS-SOCO2011	$1.11E + 01$	$1.97E + 01$	$3.08E + 01$
11(11)	PPSO	$3.94E + 00$	$1.77E + 01$	$2.17E + 01$
12(12)	DYYPO	$5.96E - 01$	$1.76E + 01$	$1.82E + 01$
13(13)	TLBO-FL	$2.89E - 02$	$1.68E + 01$	$1.68E + 01$

Tab. 5.7 CEC17 – Score – EBOwithCMAR

Tab. 5.8 $\rm CEC17-Score-jSO$

Rank	Algorithm	Score 1	Score 2	Score
1(1)	EBOwithCMAR	$4.89E + 01$	$5.00E + 01$	$9.89E + 01$
2(2)	iSO	$5.00E + 01$	$4.76E + 01$	$9.76E + 01$
3(4)	LSHADE SPACMA	$4.60E + 01$	$4.87E + 01$	$9.47E + 01$
4(3)	$LSHADE-enEpSin$	$4.71E + 01$	$4.35E + 01$	$9.06E + 01$
5(5)	DES	$4.55E + 01$	$4.33E + 01$	$8.88E + 01$
6(6)	MM OED	$4.55E + 01$	$3.81E + 01$	$8.36E + 01$
7(7)	IDEbestNsize	$2.96E + 01$	$2.76E + 01$	$5.72E + 01$
8(9)	RB-IPOP-CMA-ES	$3.76E + 00$	$3.38E + 01$	$3.76E + 01$
9(8)	$MOS-CEC2013$	$1.88E + 01$	$1.82E + 01$	$3.70E + 01$
10(10)	MOS-SOCO2011	$1.10E + 01$	$2.01E + 01$	$3.11E + 01$
11(11)	PPSO	$3.89E + 00$	$1.81E + 01$	$2.20E + 01$
12(12)	DYYPO	5.88E-01	$1.79E + 01$	$1.85E + 01$
13(13)	TLBO-FL	$2.84E - 02$	$1.71E + 01$	$1.72E + 01$

Rank	Algorithm	Score 1	Score 2	Score
1(3)	$LSHADE-enEpSin$	$5.00E + 01$	$5.00E + 01$	$1.00E + 02$
2(1)	EBOwithCMAR	$4.73E + 01$	$4.92E + 01$	$9.66E + 01$
3(2)	iSO	$4.73E + 01$	$4.62E + 01$	$9.35E + 01$
4(4)	LSHADE SPACMA	$4.45E + 01$	$4.81E + 01$	$9.26E + 01$
5(5)	DES	$4.40E + 01$	$4.34E + 01$	$8.74E + 01$
6(6)	MM OED	$4.40E + 01$	$3.79E + 01$	$8.19E + 01$
7(7)	IDEbestNsize	$2.86E + 01$	$2.75E + 01$	$5.61E + 01$
8 (9)	RB-IPOP-CMA-ES	$3.63E + 00$	$3.37E + 01$	$3.73E + 01$
9(8)	$MOS-CEC2013$	$1.81E + 01$	$1.83E + 01$	$3.64E + 01$
10(10)	MOS-SOCO2011	$1.06E + 01$	$2.02E + 01$	$3.09E + 01$
11(11)	PPSO	$3.76E + 00$	$1.82E + 01$	$2.20E + 01$
12(12)	DYYPO	$5.68E - 01$	$1.80E + 01$	$1.86E + 01$
13(13)	TLBO-FL	$2.75E - 02$	$1.72E + 01$	$1.73E + 01$

Tab. 5.9 CEC17 – Score – LSHADE-cnEpSin

Tab. 5.10 CEC20 – Score – Default BCM

Rank	Algorithm	Score 1	Score 2	Score
	i2020	$5.00E + 01$	$4.56E + 01$	$9.56E + 01$
$\mathcal{D}_{\mathcal{L}}$	IMODE.	$2.14E + 01$	$5.00E + 01$	$7.14E + 01$
3	AGSK	$2.31E + 01$	$4.56E + 01$	$6.88E + 01$

Tab. 5.11 CEC20 – Score – IMODE

Rank	Algorithm	Score 1	Score 2	Score
1(1)	i2020	$5.00E + 01$	$4.51E + 01$	$9.51E + 01$
2(2)	IMODE	$2.34E + 01$	$5.00E + 01$	$7.34E + 01$
3(3)	AGSK	$2.30E + 01$	$4.47E + 01$	$6.77E + 01$

Tab. 5.12 CEC20 – Score – AGSK

Rank	Algorithm	Score 1	Score 2	Score
1(1)	i2020	$5.00E + 01$	$4.52E + 01$	$9.52E + 01$
2(2)	IMODE	$2.14E + 01$	$5.00E + 01$	$7.14E + 01$
3(3)	AGSK	$2.25E + 01$	$4.43E + 01$	$6.68E + 01$

Tab. 5.13 CEC20 – Score – j2020

Algorithm	Score 1	Score 2	Score
$LSHADE-enEpSin$	$5.00E + 01$	$5.00E + 01$	$1.00E + 02$
EBOwithCMAR	$4.77E + 01$	$5.00E + 01$	$9.76E + 01$
iSO	$4.84E + 01$	$4.58E + 01$	$9.42E + 01$
LSHADE SPACMA	$4.45E + 01$	$4.78E + 01$	$9.23E + 01$
DES	$4.40E + 01$	$4.30E + 01$	$8.70E + 01$
MM OED	$4.40E + 01$	$3.81E + 01$	$8.21E + 01$
IDEbestNsize	$2.86E + 01$	$2.76E + 01$	$5.62E + 01$
RB-IPOP-CMA-ES	$3.63E + 00$	$3.37E + 01$	$3.73E + 01$
$MOS-CEC2013$	$1.81E + 01$	$1.82E + 01$	$3.64E + 01$
MOS-SOCO2011	$1.06E + 01$	$2.02E + 01$	$3.08E + 01$
PPSO	$3.76E + 00$	$1.82E + 01$	$2.20E + 01$
DYYPO	$5.68E - 01$	$1.80E + 01$	$1.86E + 01$
TLBO-FL	$2.75E - 02$	$1.72E + 01$	$1.73E + 01$

Tab. 5.14 CEC17 – Score

Tab. 5.15 CEC20 – Score

Rank	Algorithm	Score 1	Score 2	Score
1(1)	i2020	$5.00E + 01$	$4.19E + 01$	$9.19E + 01$
2(2)	IMODE.	$3.14E + 01$	$5.00E + 01$	$8.14E + 01$
3(3)	AGSK	$2.96E + 01$	$4.53E + 01$	$7.49E + 01$

For the CEC17, tables [5.7,](#page-55-0) [5.8,](#page-55-1) and [5.9](#page-56-1) represent the situations when only one algorithm selects its best variant of the BCM. If the rank is changed against Table [5.6,](#page-54-0) the original rank is shown in parentheses. The most noticeable difference is in Table [5.9,](#page-56-1) where the LSHADE-cnEpSin obtained the first rank. Table [5.14](#page-57-0) then contains the ranks accomplished if all three algorithms had used the bestperforming variant of the BCM, and again, the LSHADE-cnEpSin would have achieved the first position.

For the CEC20, the process is the same as for CEC17. The results are presented in Table [5.10](#page-56-0) - [5.15](#page-57-1) and no change in the algorithms order was observed.

5.4.7 Key findings

The motivation behind this contribution was to establish if the BCM can influence the algorithm performance from the competition results point of view. Thus, raise awareness about the need for careful selection of the BCM, similar to other hyperparameters of the metaheuristic algorithms. The presented results confirm that ill-selected BCM can negatively influence the algorithm's overall performance.

While the boundary control methods (BCM) are often an overlooked part of the experiment design in metaheuristics benchmarking, the paper aimed to highlight the importance of understanding the BCM as a necessary input for results reproducibility. Further, the attention was to the possibility of performance improvement by the use of alternate boundary control methods, as presented in the results of CEC17 benchmark participants, where:

- The LSHADE-cnEpSin algorithm would have won the CEC17 competition, if it did employ the random boundary control method.
- According to the scores as defined by the CEC17 benchmark, none of the three tested algorithms achieved the best results with the original BCM.

In other words, it was possible to improve the results of any of these algorithms by using a different BCM.

• Only five of the 12 participants reported on the employed BCM in the papers (albeit in two cases only partially, see Table [5.1\)](#page-44-0). Moreover, the results of the two best-performing algorithms would be irreproducible without a detailed study of the source code.

The results of the second experiment, using the CEC20 benchmark seem to be less conclusive, as only three scenarios show statistically significant differences in performance (see Table [5.4\)](#page-51-0). However, the scoring (Tables [5.5](#page-49-0) d), [5.5](#page-49-0) e) and [5.5](#page-49-0) f)) seems to indicate not only a difference in performance, but a possible benefit of using other than default BCM.

To conclude, the findings highlight a significant gap in the reproducibility of results among competition entries, primarily due to the omission of information about the utilized BCM. This oversight not only hampers the reproducibility of results but also overlooks potential performance enhancements that could be achieved by focusing on BCMs. Moreover, it has been observed that algorithms, particularly from the Differential Evolution (DE) family, implemented in Matlab, often rely on the same or similar libraries for BCM. These libraries commonly include the implementation of the halving BCM, likely influencing researchers' preference for its use due to its ready availability.

5.5 Exploring the Frequency of BCMs Activation

This study delves into the relationship between the frequency of BCM activation and various problem characteristics, such as dimensionality and fitness landscape, analyzing each dimension separately. The focus was on evaluating the top three algorithms from the CEC20 competition (AGSK, IMODE, and j2020) using the competition's benchmark set.

5.5.1 Experimental setup

The activation frequency of BCMs was assessed for each function in the benchmark set across the top three performing algorithms. The number of function evaluations (FEs) and the population size were standardized according to competition rules to align with original benchmarking conditions. Each experimental setting was conducted 30 times to ensure the reliability of the results, and the average number of BCM activations was calculated for each problem dimension. Dimension sizes of 5 and 10 were specifically examined.

To offer a comprehensive understanding of BCM activation patterns, activations were observed in each problem dimension separately. Through this analytical approach, insights into the intricacies of BCM activation frequency and its relationship with the problem's dimensionality were gained. Furthermore, it was examined whether the BCM activation frequency differed significantly across the functions in the benchmark set, yielding valuable information for algorithm designers and researchers.

The results are presented using stacked graphs, a highly effective visualization technique that provides several advantages for displaying and interpreting data. Stacked graphs are utilized for a clear and concise representation of multiple datasets within a single, unified plot. In the context of this research, stacked graphs are employed to effectively illustrate the activation frequency of various BCMs in relation to problem dimensionality and fitness landscape. Additionally, the same technique is used to visualize differences in BCM activation rate among the distinct problem dimensions.

Figures Fig. [5.7](#page-62-0) and Fig. [5.8](#page-63-0) display stacked graphs of the average number of BCM activations for three different algorithms (IMODE, AGSK, and j2020) across six different BCMs. In each figure, each column represents an algorithm, with BCMs stacked on top of each other to form a bar chart. The x-axis displays the six different BCMs, and the y-axis indicates the average number of activations (over 30 runs), for each algorithm-dimension combination. It is important

to note that the values for each algorithm in the stacked graph are represented as the sum of BCM activations in different dimensions. Different colors in each column represent the different algorithms.

Figures Fig. [5.9](#page-64-0) and Fig. [5.10](#page-65-0) show different stacked graphs; these depict a particular BCM activation on a selected test function, dimension size, and algorithm. The x-axis represents the function evaluations (FEs) of the algorithm, and the y-axis shows the average number of activations (over 30 runs) for each problem dimension.

From the analysis of the first group of graphs (Fig. [5.7](#page-62-0) and Fig. [5.8\)](#page-63-0), it is concluded that the AGSK algorithm contributes to BCM activation the most across all test functions and both dimension sizes. It is also concluded that the particular BCM affects the AGSK algorithm the most. Moreover, the number of BCM activations for algorithms j2020 and IMODE is observed to remain fairly consistent across all BCM variants.

From Figures Fig. [5.9](#page-64-0) and Fig. [5.10,](#page-65-0) it is evident that the number of BCM activations is dependent on the problem - specifically the test function. The number of BCM activations is almost equally distributed over the dimensions for f_2 (Fig. [5.9](#page-64-0) (b)), whereas for test function f_1 (Fig. 5.9 (c)), the distribution is uneven, with the number of BCM activations for dimension two prevailing over the averaged algorithm run.

All figures are included in lower quality at the end of this thesis as supplementary data. For a more detailed view, high-resolution versions of the same figures are accessible from a designated webpage $¹$ together with codes of the examined</sup> algorithms and their results.

 $^{\rm 1)}$ <https://go.fai.utb.cz/2023workshop>

Fig. 5.7 Total BCM activation for $dim = 5$. [\[116\]](#page-86-1)

Fig. 5.8 Total BCM activation for $dim = 10$. [\[116\]](#page-86-1)

Fig. 5.9 BCM activation over the run of the algorithm. [\[116\]](#page-86-1)

Fig. 5.10 BCM activation over the run of the algorithm. [\[116\]](#page-86-1)

5.5.2 Key findings

Based on the findings that reveal variations in BCM activation rates across different algorithms, test problems (fitness landscapes), and problem dimensions, as well as differences in activation rates for each BCM on the same problem, several important conclusions are drawn:

Algorithm-specific characteristics: Differences in BCM activation rates among different algorithms are observed, indicating unique characteristics for each algorithm that influence how boundary constraints are handled. This underscores the importance of selecting appropriate algorithms for specific problem types and suggests potential improvements through a better understanding of BCM behavior.

Problem-dependent activation rates: Variations in BCM activation rates across different fitness landscapes suggest that the effectiveness of BCMs is strongly dependent on the characteristics of the test problems. This necessitates careful consideration of problem-specific properties when designing or selecting BCMs to ensure optimal performance.

Dimensionality impact: The variability of BCM activation rates among dimensions of the same problem highlights the influence of problem dimensionality on the complexity of boundary constraints. This emphasizes the need to consider dimensionality's impact on BCM activation patterns when designing or adapting algorithms for high-dimensional optimization challenges.

Tailoring BCMs for improved performance: Observed differences in activation rates for each BCM on the same problem suggest that a universal solution for boundary control does not exist. By understanding these variations and identifying the most effective BCM for a given problem or algorithm, researchers and practitioners can tailor BCM implementation to enhance performance in optimization tasks.

6 THE CONTRIBUTION TO SCIENCE AND PRAC-TICE

Metaheuristic-based optimization methods are currently enjoying immense popularity. Alongside this growing popularity, the volume of research articles on this subject is also expanding, with continuous development and modification of new and existing algorithms respectively. A crucial aspect of this development process is the meticulous control of parameters that govern the behavior of the algorithm. Given that the primary focus is often on algorithm performance, selecting optimal control parameters is critical. The term "control parameter" encompasses a range of variables, including information storage systems and the selection of methods for managing critical states.

The BCMs are categorized as procedures for handling these critical states, as they are applied when a trial solution falls into an infeasible space. Preliminary studies indicate that each BCM may affect algorithm performance differently, necessitating careful consideration during the selection process. An ill-advised selection of BCMs can degrade the performance of the metaheuristic algorithm or alter other behaviors, potentially compromising specific desired characteristics of the algorithm.

Research articles that propose new metaheuristic algorithms or modifications to existing ones often overlook BCMs. Without specific details on the BCMs used by the original authors, subsequent implementations of such algorithms may be imprecise, leading to variations in effectiveness when solving specific tasks.

Addressing the research gap described above, which focuses on the often overlooked BCMs, constitutes the main part of this thesis. The first goal of this work is to raise awareness within the scientific community about the importance of BCMs, demonstrated by the presented results which prove their real impact on algorithm performance. This impact is evident not only in basic metaheuristic algorithms but also in state-of-the-art variants that have participated in benchmark competitions.

The second goal targets algorithm designers, who are urged to pay careful attention to providing a detailed description of the algorithm and its setup. This is crucial for the reproducibility of results and the effective evaluation of algorithm performance across different implementations.

The work highlights the significance of BCMs in the development and benchmarking of metaheuristic algorithms, and BCMs should also be important components in the automatic design or configuration of algorithms. It is imperative that these components are not merely mentioned as afterthoughts but are integrated into the core design and reporting of algorithmic research to ensure accuracy and replicability in scientific studies.

7 GOAL FULFILLMENT

This section outlines the steps implemented to achieve the dissertation goal, which were established as follows:

- $\sqrt{\ }$ Survey the current state of boundary control methods (BCMs) used in evolutionary algorithms: The current state of BCMs was extensively reviewed in Section [4.1,](#page-27-0) where a comprehensive survey of the literature was conducted not only to establish a foundational understanding of BCM applications in evolutionary algorithms but also to investigate which BCMs are being utilized. Additionally, this review explored whether other researchers, particularly in the context of algorithm design and benchmarking, are addressing these issues.
- $\sqrt{\ }$ Investigate the influence of various BCMs on the performance of selected evolutionary algorithms: Initial studies, detailed in Sections [5.1,](#page-35-0) [5.2](#page-36-0) and [5.3,](#page-40-0) investigated the impact of BCMs on both basic and more advanced variants of various algorithms such as PSO, FA, and SOMA. These investigations laid the groundwork for subsequent experiments aimed at assessing the influence of BCMs on state-of-the-art algorithms.
- \checkmark Conduct experiments evaluating the impact of BCMs on the performance of state-of-the-art algorithms and results of competitive benchmarking: The experiments were carried out as described in Section [5.4](#page-43-0) and Section [5.5,](#page-59-0) which tested the efficacy and the influence of BCMs on state-of-the-art algorithms using modern competitive benchmarks. The results from these experiments corroborated the preliminary findings from the initial studies, confirming the significant impact of BCMs on algorithm performance.
- $\sqrt{\ }$ Based on the experimental results, draw conclusions and recommendations for good practices in benchmarking: The conclusions and recommendations were presented, underscoring the significant influence of BCMs on the performance of metaheuristic algorithms. As evidenced by the experimental results in Section [5.4,](#page-43-0) appropriately chosen

BCMs significantly altered the competitive ranking of the algorithms. The analysis of competition algorithms and the survey of state-of-the-art literature suggest that BCMs should be considered an integral part of the hyperparameter analysis of any algorithm design. Authors should always specify which BCM implementation was chosen to ensure fair comparison and reproducibility.

These findings underscore the importance of BCMs in the design and analysis of metaheuristic algorithms, advocating for their consistent inclusion in algorithmic research and documentation.

8 CONCLUSION

This dissertation provides a comprehensive treatise on the use of Boundary Control Methods (BCMs) in metaheuristic algorithms. The initial sections introduce the fundamental concepts and relationships among these fields, with a specific focus on the role and impact of BCMs in contemporary EA trends as outlined in subsection [3.6.](#page-24-0)

Following the introduction, the dissertation delineates the proposed goals and the methodologies employed to achieve them. A detailed examination of BCMs and a state-of-the-art overview are presented, where five representative BCMs are selected based on the literature review. The methodology and preliminary results, previously published at international conferences by the Author, are described in subsequent sections. After these preliminary results, further experiments were conducted to evaluate the impact of BCMs on the performance of state-of-the-art algorithms and the results of competitive benchmarking. Additionally, an experiment was included to investigate the frequency of BCM usage among state-of-the-art algorithms in the CEC20 benchmark.

While BCMs are often an overlooked part of experiment design in metaheuristics benchmarking, this dissertation highlights the importance of understanding BCMs as a necessary input for results reproducibility and potential performance improvement. The experimental findings from the CEC17 benchmark participants clearly demonstrate this. Notably, the LSHADE-cnEpSin algorithm could have won the CEC17 competition if it had employed the *random* BCM. Additionally, it was observed that none of the three tested algorithms achieved the best results with their original BCMs, indicating that performance improvements were possible through alternative BCMs. However, only five of the 12 CEC17 participants provided details on their BCM practices, with two of these reports being incomplete, which significantly impairs the reproducibility of their experiments. These findings underscore the significant influence of BCMs on the performance of metaheuristic algorithms, as evidenced in Section 5.4, where appropriately chosen BCMs significantly altered the competitive ranking of the
algorithms. The analysis of competition algorithms and the survey of state-ofthe-art literature suggest that BCMs should be considered an integral part of the hyperparameter analysis of any algorithm design. Authors should always specify which BCM implementation was chosen to ensure fair comparison and reproducibility, advocating for their consistent inclusion in algorithmic research and documentation.

This dissertation has conclusively demonstrated that BCMs are not merely supplementary components but are integral to the effective design, analysis, and application of metaheuristic algorithms. Their role should be carefully considered and integrated into future research and practice in the field of optimization. The importance of future research on BCMs lies in their universal applicability and profound impact across the entire field of metaheuristic optimizers, particularly in bound-constrained scenarios. Such research is crucial for advancing our understanding and implementation of these methods, ensuring they contribute significantly to the robustness and efficacy of optimization solutions.

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CURRICULUM VITAE

LIST OF APPENDICES

APPENDIX A: Total BCM activation frequency results

APPENDIX B: BCM Activation Dynamics

APPENDIX A: TOTAL BCM ACTIVATION FREQUENCY RESULTS

 $dim = 5, f_1$ $dim = 5, f_2$ $dim = 5, f_3$

$$
dim=5, f_4
$$

 $\bar{1}$ $\frac{80}{2}$ activation

 $dim = 5, f_9$ $dim = 5, f_{10}$ $dim = 10, f_1$ $dim = 10, f_2$

 $dim = 10, f_3$ $dim = 10, f_4$ $dim = 10, f_5$ $dim = 10, f_6$

APPENDIX B: BCM ACTIVATION DYNAMICS

Algorithm j2020, BCM Default

Algorithm j2020, BCM Clipping

Algorithm j2020, BCM Random

Algorithm j2020, BCM Periodic

Algorithm j2020, BCM Reflection

Algorithm j2020, BCM Halving

Algorithm IMODE, BCM Default

Algorithm IMODE, BCM Clipping

Algorithm IMODE, BCM Random

Algorithm IMODE, BCM Periodic

Algorithm IMODE, BCM Reflection

Algorithm IMODE, BCM Halving

Algorithm AGSK, BCM Default

Algorithm AGSK, BCM Clipping

Algorithm AGSK, BCM Random

Algorithm AGSK, BCM Periodic

Algorithm AGSK, BCM Halving

