An Improved Estimation of Distribution Algorithm for Mixed-Integer Nonlinear Programming Problems: $EDAII_{mv}$

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Abstract. In a mixed-integer nonlinear programming problem, integer restrictions divide the feasible region into discontinuous feasible parts with different sizes. Meta-heuristic optimization algorithms quickly lose diversity in such scenarios and get trapped in local optima. In this work, we propose an Estimation of Distribution Algorithm (EDA) with two modifications from its previous version (EDA_{mv}) . The first modification consists in establishing the exploration and exploitation components for the histogram of discrete variables, aimed at improving the performance of the algorithm The second modification is during the evolution. a repulsion operator to overcome the population stagnation in discontinuous parts, so as continuing the search for possible solutions in other regions. From a comparative study on 16 test problems, the individual contribution of each modification was verified. According to statistical test results, the new proposal shows a significantly better performance than the other competitors tested.

Keywords. Estimation of distribution algorithm, integer restriction handling, mixed integer nonlinear programming.

1 Introduction

Many optimization problems, especially in the field of engineering, have variables that cannot take every value in a continuous space. Instead, such variables can only take integer values, or discrete values in the general sense. Integer variables are commonly used to define elements of the same class, e.g., worker assignment, car control with gear change, multi-stage mill design, selection of standardized elements, etc. Nonlinear problems where continuous, integer, and discrete variables coexist are known as Mixed-Integer Nonlinear Programming (MINLPs) problems [10]. In general, a MINLP problem can be defined by (1) – (6):

$$\min f(\mathbf{x}, \mathbf{y}),$$
 (1)

s.t.
$$g_i(\mathbf{x}, \mathbf{y}) \le 0, i = 1, ..., n_i,$$
 (2)

$$h_j(\mathbf{x}, \mathbf{y}) = 0, \ j = 1, ..., n_j,$$
 (3)

$$x_k^L \le x_k \le x_k^U, \ k = 1, ..., n_k,$$
 (4)

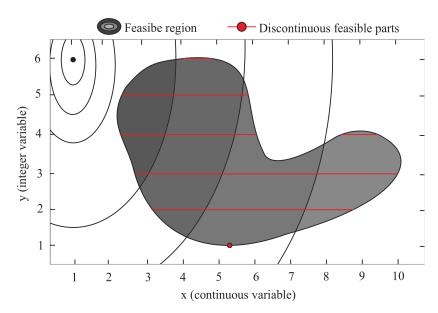


Fig. 1. MINLP problem example, where the shaded area represents the feasible region defined by the constraints, and the red lines are the discontinuous feasible parts that also satisfy the integer restrictions

$$y_q^L \le y_q \le y_q^U : \text{integer}, \ q = 1, ..., n_q,$$
 (5)

$$[\mathbf{x}, \mathbf{y}] \in \eta,$$
 (6)

where $f(\mathbf{x}, \mathbf{y})$ is the objective function, \mathbf{x} is a vector of continuous decision variables, \mathbf{y} is a vector of integer decision variables, x_k^L and x_k^U are the lower and upper bounds of x_k , respectively, y_q^L and y_q^U are the lower and upper bounds of y_q , respectively, η is the decision variable space, $g_i(\mathbf{x}, \mathbf{y})$ is the ith inequality constraint, and $h_j(\mathbf{x}, \mathbf{y})$ is the jth equality constraint.

In a MINLP problem, the integer restrictions divide the feasible region into discontinuous feasible parts with different sizes. Fig. 1 shows a MINLP problem, where \boldsymbol{x} is a continuous variable, and \boldsymbol{y} is an integer variable.

The shaded area is the feasible region defined by the constraints, and the red lines are the discontinuous feasible parts that also satisfy the integer restrictions.

In recent years, meta-heuristic optimization algorithm have gained popularity over classical MINLP techniques.

Different extensions of genetic algorithms [2], particle swarm optimization [4, 16], differential

evolution [1, 5], ant colony optimization [13], harmony search [3], estimation of distribution algorithm [15], aimed at solving MINLP problems have been proposed.

The most significant advantage of these algorithms is their robustness regarding the function properties, such as non-convexity or discontinuities [12].

The classical MINLP techniques (like branch and bound, cutting planes, outer approximation) generally require prior convexification and relaxation operations, which are not always possible [11].

On the other side, when the population of meta-heuristic optimization algorithm converges to a discontinuous feasible part, it quickly loses diversity, and the exploration is reduced, with no possibility of jumping out to another discontinuous feasible part. Compared to larger discontinuous parts, it is difficult to find feasible solutions in the smaller parts. If the best solutions are located in small parts, then the population might converge to the wrong solutions.

Only a few recent works focused on MINLP problems consider the drawbacks described above. In [7], a multiobjective differential evolution is proposed.

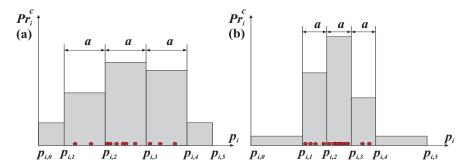


Fig. 2. Search progress of the AWH model for W = 3. (a) first generations, (b) later generations

This strategy gives equal priority to integer conditions and quality of the solution, and the population converges to good regions regarding both criteria.

In [6], the authors propose a cutting strategy that penalizes non-promising solutions, which means that non-promising parts are progressively discarded.

In addition, they propose a repulsion strategy that penalizes the discontinuous parts where the population is trapped, in order to search better solutions in other regions.

More recently, in [9] the Estimation of Distribution Algorithm for Mixed-Variable Newsvendor problem (EDA_{mvn}) [15] is improved and proposed to MINLP problems.

The new proposal (EDA_{mv}) uses the ε -constrained method to explore the smaller discontinuous feasible parts from infeasible contours. Also, the hybridization with a mutation operator is proposed.

In this work, we propose an algorithm, $EDAII_{mv}$, with two modifications from the original EDA_{mv} .

The first modification consists in establishing the exploration and exploitation components for the histogram of discrete variables, using the balance between both terms to improve the performance of the algorithm during the evolution.

The second modification is a repulsion operator to overcome the population stagnation in discontinuous parts, and continue the search for possible solutions in other regions.

Through a comparative analysis, the individual contribution of each modification to the algorithm

performance was verified. The performance of $EDAII_{mv}$ is significantly higher than those of the compared algorithms.

2 Estimation of Distribution Algorithm

 EDA_{mv} is an improved version of EDA_{mvn} , originally proposed in [15]. It uses an Adaptive-Width Histogram (AWH) model for handling continuous variables, and an ε -linked Learning-Based Histogram (LBH ε) model for handling discrete variables.

New variable values are generated from statistical sampling. In the case of continuous variables, statistical sampling is hybridized with a mutation operator.

The replacement mechanism to get the next population is carried out through parent-offspring competition using the ε -constrained method.

2.1 Adaptive-width Histogram Model

The AWH model promotes promising regions by assigning them high probabilities, while in the other regions very low probabilities are assigned. One AWH is developed for each decision variable independently.

The search space $[a_i,b_i]$ of the ith variable x_i is divided into (W+2) bins (regions), to define the probabilities Pr_i^c for the AWH model.

Points $[p_{i,0},p_{i,1},...,p_{i,w+1},p_{i,w+2}]$ define the width of the bins shown in Fig. 2, where $p_{i,0}=a_i$ and $p_{i,w+2}=b_i$ (a_i and b_i are the lower and upper bounds of x_i , respectively).

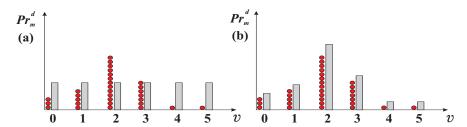


Fig. 3. LBH ε model for v = 6. (a) $\varepsilon > \varepsilon_p$ equal probability for all available integer values, (b) $\varepsilon \le \varepsilon_p$ considering population distribution

The total number of bins is (W+2) although the input parameter for EDA_{mv} is W, since the algorithm creates two more bins: one between the lower boundary a_i and the point $p_{i,1}$, and another one between the point $p_{i,w+1}$ and the upper boundary b_i (unpromising regions).

By assuming that $x_{i,\min}^1$ and $x_{i,\min}^2$ are the smallest and the second smallest existing values of variable x_i , respectively, and $x_{i,\max}^1$ and $x_{i,\max}^2$ are the highest and the second highest existing values of variable x_i , respectively, then points $p_{i,1}$ and $p_{i,w+1}$ are defined as in (7) and (8):

$$p_{i,1} = \max\{x_{i,\min}^1 - 0.5(x_{i,\min}^2 - x_{i,\min}^1), p_{i,0}\},$$
 (7)

$$p_{i,w+1} = \min\{x_{i,\max}^1 + 0.5(x_{i,\max}^1 - x_{i,\max}^2), p_{i,w+2}\},$$
 (8)

The W bins of the promising areas are located in the range $[p_{i,1}, p_{i,w+1}]$, and have the same width a, given by (9):

$$a = \frac{(p_{i,w+1} - p_{i,1})}{W}. (9)$$

Let $A_{i,j}$ be the count of individuals for the ith variable located in the jth bin. As can be seen in Fig. 2, the end bins do not contain solutions (unpromising regions), then $A_{i,1} = A_{i,W+2} = 0$.

However, a small value will be assigned through the parameter e_b , to avoid premature convergence. $A_{i,j}$ is obtained by (10):

$$A_{i,j} = \begin{cases} A_{i,j}, & \text{if } 2 \leq j \leq (W+1), \\ e_b, & \text{if } j=1, (W+2), \text{ and } p_{i,j} > p_{i,j-1}, \\ 0, & \text{if } j=1, (W+2), \text{ and } p_{i,j} = p_{i,j-1}. \end{cases} \tag{10}$$

The first case in (10) is the count of bins of promising regions $[p_{i,1}, p_{i,w+1}]$. The second case corresponds to unpromising regions with e_b value.

The third case assigns zero to the end bins with empty range. The probability of the ith variable in the jth bin is obtained by (11):

$$Pr_{i,j}^{c} = \frac{A_{i,j}}{\sum_{k=1}^{W+2} A_{i,k}}.$$
 (11)

2.2 Learning-based Histogram Model Linked with ε -constrained

The LBH ε model is used for handling integer variables. It is a link between the LBH model and the ε -constrained method.

The aim is to maintain an equal probability for all available integer values until ε reaches a predefined value ε_p , as is shown in Fig. 3 (a).

When ε reaches ε_p , the LBH model begins the learning process, i.e., considering the information of the population distribution to update the probability, as shown in Fig. 3 (b).

If the ε -constrained method has been effective, for values of ε sufficiently small, the solutions must be close to those parts of the feasible region with promising objective function values.

Therefore, if the histogram begins the learning process at that point, it has a better chance of converging to good solutions.

Considering that the variable y_m has v available integer values, with $v \in \{L_m, L_m+1, L_m+2, ..., U_m\}$, the probability of the nth available value of v is defined by (12):

$$Pr_{m,v}^{d}(t) = \begin{cases} \frac{1}{(U_m - L_m) + 1}, & \text{if } \varepsilon > \varepsilon_p, \\ (1 - \gamma) \cdot Pr_{m,v}^{d}(t - 1) + \frac{\gamma \text{ Count}_v}{N}, & \text{if } \varepsilon \leq \varepsilon_p, \end{cases}$$
 (12)

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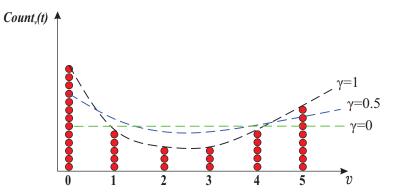


Fig. 4. LBH ε model, γ =0 random exploration, γ =0.5 middle consideration of population distribution, γ =1 total consideration of population distribution

where N is the population size, t is the current generation, γ is the population learning rate, and Count_v is the number of individuals with the $n\mathrm{th}$ available value of v.

Let t_{max} be the maximum number of generations, and γ a dynamic parameter defined by (13):

$$\gamma = \frac{t}{t_{max}}. (13)$$

Therefore, as the number of generations advances, γ gradually increases as well, which implies an accelerated learning process, i.e., the model uses more information of the current population distribution.

2.3 Sampling

After the histograms have been developed, the offspring is obtained by sampling the models.

In case of a continuous variable x_i , a bin j is firstly selected according to a randomly generated probability, then x_i is uniformly sampled from the points that limit the bin selected $[p_{i,j-1},p_{i,j})$.

For a discrete variable y_m , an available value of $v \in \{L_m, U_m\}$ is selected by a randomly generated probability.

2.4 Hybridization with a Mutation Operator

The mutation operation is added to generate the real variables. The vector of real variables \mathbf{x} of each offspring is generated by mutation or by sampling taking into account the predefined mutation probability r_M , i.e. if this probability is satisfied for a solution vector, its real variables are computed as shown in (14) and (15):

$$x_{k i}^{g+1} = x_{best i}^g + \beta \cdot (x_{best i}^g - x_{k i}^g),$$
 (14)

$$\beta = \beta_{\min} + \operatorname{rand}_{k,i} \cdot (\beta_{\max} - \beta_{\min}), \tag{15}$$

where $k,\ i$ are the index of the current solution vector and current variable, respectively, g is the current generation, $x_{{\rm best},i}^g$ is the ith variable of the best solution vector found so far, ${\rm rand}_{k,i}$ is a random number between 0 and 1, and $\beta_{\rm min}$ and $\beta_{\rm max}$ are the lower and upper bounds of β predefined by the user, with values between 0 and 1.

In the new proposal the values of β_{\min} and β_{\max} will always be set to 0 and 1, respectively.

2.5 Constraint Handling

The replacement mechanism to get the next population is carried out through parent-offspring competition using the ε -constrained method.

The ε -constrained method was proposed by Takahama and Sakai [14] as a constraint-handling technique.

Given two function values $f(x_1)$, $f(x_2)$, and two constraint violations $\phi(x_1)$, $\phi(x_2)$ for two points x_1

Table 1. PSO_{mv} , EDA_{mv} , EDA_{mv} (I), and $EDAII_{mv}$ results

	Status	PSO _{mv}	EDA _{mv}	EDA _{mv} (I)	EDAII _{mv}
F1	FR	100	100	100	100
	SR	0	100	0	100
	Ave ± Std Desv	17.000±0.000 +	13.000±0.000 ≈	17.000±0.000 +	13.000±0.000
F2	FR	100	100	100	100
	SR	100	100	100	100
	Ave ± Std Desv	1.000±0.000 \approx	1.000±0.000 \approx	1.000±0.000 \approx	1.000±0.000
F3	FR	100	100	100	100
	SR	24	100	76	100
	Ave ± Std Desv	-3.879±0.217 +	-4.000±0.000 $pprox$	-3.880±0.218 +	-4.000±0.000
F4	FR	100	100	100	100
	SR	100	100	100	100
	Ave ± Std Desv	-6.000±0.000 $pprox$	-6.000±0.000 $pprox$	-6.000±0.000 $pprox$	-6.000±0.000
F5	FR	100	100	100	100
	SR	0	100	76	100
	Ave ± Std Desv	1.240±0.000 +	0.250±0.000 ≈	0.488±0.432 +	0.250±0.000
	FR	100	100	100	100
F6	SR	100	100	100	100
	Ave ± Std Desv	-6,783.582±0.000 ≈	-6,783.582±0.000 ≈	-6,783.582±0.000 ≈	-6,783.582±0.000
F7	FR	96	100	100	100
	SR	0	24	28	36
	Ave ± Std Desv	NA +	0.895±0.235 +	0.725±0.361 +	0.642±0.359
F8	FR	100	92	100	100
	SR	0	0	0	0
	Ave ± Std Desv	7,222.847±94.800 -	NA +	7,971.856±518.086 ≈	7,986.723±906.139
F9	FR	100	88	100	100
	SR	16	0	0	0
	Ave ± Std Desv	7,284.444±283.224 -	NA +	8,305.496±742.746 ≈	8,391.061±854.267
F10	FR	100	64	96	100
	SR	64	0	0	0
	Ave ± Std Desv	7,337.332±277.610 -	NA +	NA +	8,086.671±641.101
F11	FR	100	100	100	100
	SR	0	0	0	0
	Ave ± Std Desv	46.280±6.601 +	40.785±5.484 +	38.119±5.378 ≈	37.822±5.334
	FR	100	100	100	100
F12	SR	0	0	0	4
	Ave ± Std Desv	90.048±17.975 +	74.500±30.941 +	51.976±20.146 ≈	56.201±23.594
F13	FR	100	100	100	100
	SR	0	0	0	4
	Ave ± Std Desv	8,956.649±7.448 ≈	8,943.236±29.864 ≈	8,955.137±31.467 ≈	8,949.792±35.701
F14	FR	100	100	100	100
	SR	0	48	60	76
	Ave ± Std Desv	8,977.707±66.813 +	8,963.673±41.007 ≈	8,954.966±10.181 ≈	8,958.233±41.392
			100	100	100
	FR	100			
F15	FR SR	0	0	0	0
F15			0 34.997±3.938 +	0 30.580±1.827 ≈	0 31.639±2.105
F15	SR	0 30.899±1.203-	34.997±3.938 +	30.580±1.827 ≈	31.639±2.105
F15	SR Ave ± Std Desv	0			
	SR Ave ± Std Desv FR	0 30.899±1.203- 100	34.997±3.938 + 100	30.580±1.827 ≈ 100	31.639±2.105 100

and x_2 , the ε -constrained method uses the ε -level comparisons described in (16):

$$(f_{1},\phi_{1}) \leq_{\varepsilon} (f_{2},\phi_{2}) = \begin{cases} f_{1} \leq f_{2}, & \text{if } \phi_{1},\phi_{2} \leq \varepsilon, \\ f_{1} \leq f_{2}, & \text{if } \phi_{1} = \phi_{2}, \\ \phi_{1} < \phi_{2}, & \text{otherwise,} \end{cases}$$
(16)

where ε -level comparisons are defined as an order relation on a pair of objective function and constraint violation values $(f(x), \phi(x))$.

This means that the candidates with a violation sum lower than ε are considered as feasible solutions and are ordered according to their fitness values.

In the case of $\varepsilon=0$, $\phi(x)$ always precedes f(x). Therefore, this method favors the approach to the feasible region by keeping slightly infeasible solutions with promising fitness values.

The ε -level decreases at each iteration G until the predefined iteration number Tc is reached, after that $\varepsilon = 0$, as indicated by (18):

$$\varepsilon(0) = \phi(x_{\theta}),\tag{17}$$

$$\varepsilon(G) = \begin{cases} \varepsilon(0)(1 - \frac{G}{T_c})^{cp}, & \text{if } 0 < G < T_c, \\ 0, & \text{otherwise,} \end{cases}$$
 (18)

where cp is a parameter to control the speed of constraint relaxation, $\varepsilon(0)$ is the initial value of ε , and x_{θ} is the top $\theta \mathrm{th}$ in an array sorted by total constraint violation ($\theta=0.2N$).

3 Proposed Method

Two modifications for EDA_{mv} are proposed.

The first proposed modification focuses on establishing a new balance between exploration and exploitation of the LBH ε model, in order to contribute to the algorithm performance during evolution.

The second modification is based on the repulsion of discontinuous parts that stagnate the population, with the aim of seeking better solutions in other discontinuous parts.

3.1 LBH ε Improvement

As described in (12), γ is the learning rate of the population. A high value of γ increases the role of the population distribution Count_v/N in obtaining the $Pr_m^d(t)$, whereas a low value mainly considers the histogram of the previous generation, $Pr_m^d(t-1)$.

However, when certain admissible values begin to prevail statistically over others, the histograms and the populations begin to be similar, so the terms of the equation (12), instead of combining different information, emphasize the same search direction and cause accelerated (and often premature) convergence.

In this work, the following LBH ε model is proposed:

$$Pr_{m,v}^{d}(t) = \begin{cases} Pe_{m,v}^{d}, & \text{if } \varepsilon > \varepsilon_{p}, \\ (1-\gamma) \cdot Pe_{m,v}^{d} + \gamma \cdot \frac{\text{Count}_{v}}{N}, & \text{if } \varepsilon \leq \varepsilon_{p}, \end{cases}$$
(19)

where Pe_m^d are equal probabilities for all v values of the mth variable, and are given by (20):

$$Pe_{m,v}^d = \frac{1}{(U_m - L_m) + 1}. (20)$$

In this model, Pe_m^d contributes to the exploration of the algorithm, while Count_v/N contributes to the exploitation on the most populated regions (promising regions).

As can be seen in Fig. 4, now for very low values of γ , the histogram will be flatter (low selection pressure).

As the value of γ increases, the histogram and selection pressure will be more consistent with the population distribution. As in the previous case, γ is a dynamic parameter defined by (13).

3.2 Repulsion

The repulsion strategy proposed in [6] consists of two steps: (i) judge whether the population is trapped into a solution, and (ii) apply a repulsion operator to the discontinuous feasible part containing the solution, and restart the population. Eq. (21) is the fail consideration to find a better solution:

$$(f_{\text{best}} - f'_{\text{best}}) \le 0 \& (g_{\text{best}} - g'_{\text{best}}) \le 0,$$
 (21)

where f_{best} and g_{best} are the objective function value and the degree of constraint violation of the best solution found so far, respectively, f_{best}' and g_{best}' are the objective function value and the degree of constraint violation of the best solution in the current generation, respectively.

If (21) is satisfied, it means that the algorithm fails to find a better solution, then the counter is incremented (ctr = ctr + 1). If (21) is not satisfied in any generation, the counter is reset (ctr = 0).

If ctr is greater than a predefined failure threshold T, the population is considered to be trapped in a solution, and the discontinuous feasible part (\mathbf{y}) containing that solution has been explored. Then the population is regenerated, and the solution is recorded in the store archive. Any population member that has a vector \mathbf{y} contained in store will be penalized with an arbitrarily large degree of constraint violation.

arepsilon-constrained method is also restarted but with a new T_c value with fewer generations, called fast generation control (T_c') . At the end of the execution, the recorded solutions should be considered to return the best solution.

4 Experimentation and Results

4.1 Benchmark Problems

Sixteen MINLP problems (F1-F16) were used to evaluate the performance of $EDAII_{mv}$. Because of the space limitation, a detailed description of the problems is not included, but it can be found in [6].

The maximum number of objective function evaluations was set at 200,000, and 25 independent runs were executed for each problem. The tolerance value for the equality constraints was set at 1.0E-04.

A run was considered as successful if: $|f(x_{\mathrm{best}}) - f(x^*)| \leq 1.0$ E-4, where x^* is the best known solution and x_{best} is the best solution provided by the algorithm.

4.2 Algorithms and Parameter Settings

 PSO_{mv} [16], EDA_{mv} [9], and $EDAII_{mv}$ were the competing algorithms in the experiment. PSO_{mv} also uses the LBH model for handling discrete variables. However, the γ is an adaptive parameter, and the LBH probability is updated using only the best half of the swarm.

To prove the individual contribution of each modification proposed, the instance with only $LBH\varepsilon$ improvement $(EDA_{mv}(I))$ was also included. The algorithms were tuned using the iRace parameter tuning tool [8]. The parameter values were as follows:

- PSO_{mv} : swarm size N=300, acceleration coefficient c=1.5299, learning rate $\gamma=0.0125$.
- EDA_{mv} : N=50, numbers of bins W=4, end bins parameter $e_b=2.3959$, control generation $T_c=3,000$, control speed parameter cp=8, link parameter $\varepsilon_p=0.2399$, and mutation parameters: $r_M=0.6$, $\beta_{\min}=0.3$, $\beta_{\max}=0.9$.
- $EDA_{mv}(I)$: N = 50, W = 3, $e_b = 2$, $T_c = 2000$, cp = 7, $\varepsilon_p = 5$, and $r_M = 0.3$.
- $EDAII_{mv}$: N=50, W=3, $e_b=2$, $T_c=2000$, cp=7, $\varepsilon_p=5$, $r_M=0.3$, failure threshold T=400, and fast control generation $T_c'=200$.

4.3 Analysis of Results

Table 1 summarizes the results of PSO_{mv} , EDA_{mv} , EDA_{mv} (I), and $EDAII_{mv}$.

These results are assessed considering the terms Feasible Rate (FR), Successful Rate (SR), Average (Ave), and Standard Deviation (Std Dev), over 25 independent runs. "NA" means that an algorithm cannot achieve 100% FR.

 EDA_{mv} (I) beats EDA_{mv} in nine problems (F7:F12, F14:F16) in at least one of the term concerned, proving that $LBH\varepsilon$ has a positive influence on the algorithm performance.

As mentioned above, the LBH model (used in EDA_{mv}) has two terms that could contain redundant information, producing an accelerated convergence.

However, for problems F1, F3, and F5, where the solutions are in small feasible parts, a slower

Computación y Sistemas, Vol. 27, No. 1, 2023, pp. 163–172 doi: 10.13053/CyS-27-1-4532

convergence of $LBH\varepsilon$ (used in $EDA_{mv}(I)$) causes that the ε -level reaches zero value when the histogram has not yet converged to the small promising part.

The repulsion strategy is very useful for this situation, since restarts the exploration in the remaining unexplored parts.

As can be seen, the implementation of repulsion strategy in $EDAII_{mv}$ improves the performance for problems F1, F3, and F5 without compromising the rest of the problems.

A Wilcoxon's rank-sum test at a 0.05 significance level was carried out between $EDAII_{mv}$ and each competitor, in order to evaluate the significant differences in the results.

In Table 1, [+], $[\approx]$ and [-] denote that $EDAII_{mv}$ is better than, worse than, and similar to its current competitor, respectively.

As shown in the final part of Table 1, the results of $EDAII_{mv}$ are significantly better than EDA_{mv} in eight problems (F7:F12, F15,F16), similar in another eight problems (F1:F6, F13, F14), and in no case EDA_{mv} surpasses the result of the new proposal.

 $EDA_{mv}(I)$ results are outperformed on five problems (F1, F3, F5, F7, F10), matched on eleven problems (F2, F4, F6, F8, F9, F11:F16), and in no case is $EDAII_{mv}$ outperformed by $EDA_{mv}(I)$.

It is clear that $EDAII_{mv}$ has significantly better results than previous variants. Analyzing the results of this sequenced implementation, it can be concluded that each proposed modification contributes to a better performance.

Regarding PSO_{mv} , the new proposal is significantly better in seven test problems (F1, F3, F5, F7, F11, F12, F14) and no difference in four problems (F2, F4, F6, F13), while PSO_{mv} outperformed $EDAII_{mv}$ in five problems (F8, F9, F10, F15, F16).

Although in general $EDAII_{mv}$ has a better performance than PSO_{mv} , the advantage of PSO_{mv} in the last mentioned problems is due to a superior diversity in the exploration.

Therefore, it is recommended in future works to focus on promoting greater diversity in $EDAII_{mv}$.

5 Conclusion and Future Work

 $EDAII_{mv}$ was proposed with two modifications regarding its previous version EDA_{mv} .

The first modification establishes a better balance between the exploration and exploitation terms in LBH ε , aimed at improving the performance of the algorithm during the evolution.

The second modification is a repulsion operator to overcome the population stagnation in discontinuous parts, and continue the search for good solutions in other regions.

Through a comparative analysis on sixteen test problems, the individual contribution of each modification to the algorithm performance was verified.

According to the Wilcoxon's rank-sum, $EDAII_{mv}$ showed significantly better performance than its previous version.

The benchmark was also used to compare the performance of the improved proposal against PSO_{mv} . Overall, $EDAII_{mv}$ has a better performance than PSO_{mv} .

However, PSO_{mv} presents an advantage in some problems due to a superior diversity in the exploration. Therefore, it is recommended in future works to focus on promoting higher diversity in the $EDAII_{mv}$.

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