# Co-evolutionary Differential Evolution with Dynamic Population Size and Adaptive Migration Strategy

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## ABSTRACT

As the performance of differential evolution (DE) is significantly affected by its mutation schemes and parameter settings when solving different problems, this paper proposes a simple yet efficient co-evolutionary DE (CEDE) to enhance the algorithm performance. The CEDE algorithm uses multiple populations to optimize the problem cooperatively, with each population using different operators and/or different parameters. Moreover, as different populations may show different performance on the same problem, we further design an efficient adaptive migration strategy (AMS) to dynamically control the population size of different populations. The CEDE algorithm is tested and compared on four benchmark functions. Experimental results demonstrate the good performance of CEDE when compared with conventional DEs using different operators and/or parameters.

#### **Categories and Subject Descriptors**

I.2.8 [Artificial Intelligence]: Problem Solving, Control Methods, and Search – *Heuristic methods*; G.1.6 [Numerical Analysis]: Optimization – *Global optimization*.

## **General Terms**

Algorithms, Performance, Reliability, Experimentation

#### Keywords

Differential evolution (DE); co-evolutionary algorithm; dynamic population size; adaptive migration strategy.

## **1. INTRODUCTION**

Differential evolution (DE) was proposed by Price and Storn as a kind of evolutionary algorithm (EA) [1]. However, the performance of DE on different problems is significantly affected by its evolutionary operators, especially the mutation operator, and parameter settings, especially the 'crossover rate' CR. Therefore, many researches have been conducted in the literature to enhance the DE performance by designing efficient mutation scheme or/and adaptively controlling the parameter settings [2]. However, using adaptive or self-adaptive strategies requires the users to understand the evolutionary process well so as to elaborately design good control strategies. In this paper, we propose to use a novel co-evolutionary DE (CEDE) with adaptive migration strategy (AMS) to address the operator choice and parameter settings problems in DE algorithm.

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## 2. CEDE WITH AMS

There are M populations in CEDE, with each population using different mutation schemes (e.g., DE/best scheme or DE/rand scheme) and different parameter settings (e.g., CR=0.1 or CR=0.9). These M populations work cooperatively and each population optimizes the problem like a conventional DE. CEDE works as the following 6 steps.

Step 1: Initialization. For each population m, define its mutation scheme  $MS_m$  and two control parameters  $F_m$  and  $CR_m$ . For each individual i in the  $m^{\text{th}}$  population, initial its position  $X_{m,i}$  to a random value within the search space and then evaluate the individual's fitness  $f(X_{m,i})$ . Find out the best individual among all the populations and its fitness is denoted as *Best*.

Step 2: Evolutionary process. In every generation, perform Step 3, Step 4, and Step 5 as follows:

Step 3: For each individual i in the m<sup>th</sup> population, perform the following operations:

Step 3.1: Perform the mutation and crossover operations on  $X_{m,i}$  based on MS<sub>m</sub>,  $F_m$ , and  $CR_m$ , so as to obtain a new position  $U_{m,i}$ .

Step 3.2: Evaluate  $f(U_{m,i})$  and compare with  $f(X_{m,i})$ . If  $f(U_{m,i})$  is better than  $f(X_{m,i})$ , then  $X_{m,i}$  is set to  $U_{m,i}$ . Otherwise, keeps  $X_{m,i}$ .

Step 4: Find out the best individual among all the population and update the value of *Best*.

Step 5: Perform AMS among all the *M* populations.

AMS calculates the mean fitness value of each population and sorts the *M* populations according to the obtained mean fitness values from good to poor. All the poor populations will randomly migrate one individual to the good populations. For example, all the populations from rank 2 to rank *M* randomly migrate one individual to the 1<sup>st</sup> rank population, all the populations from rank 3 to rank *M* randomly migrate one individual to the 2<sup>nd</sup> rank population, and so on. However, AMS uses a parameter  $p_m$  as the *'migration rate'* to control the migration. Before the migration, a random real number *r* is first generated in the range [0, 1], if *r* is smaller than  $p_m$ , then the migration occurs. Otherwise, the migration does not occur. The  $p_m$  is non-linear time varying depended on evolutionary process as:

$$p_m = 0.01 + 0.99 \frac{(\exp(\frac{10g}{G}) - 1)}{(\exp(10) - 1)} \tag{1}$$

where g and G are the current generation and the maximal generation respectively.

Step 6: Termination check. If the termination condition is met, then print the value of *Best* and CEDE terminates. Otherwise, CEDE goes to Step 2 for the next generation.

#### **3. EXPERIMENTAL STUDIES**

Four functions as listed in Table 1 are used for experiments.  $f_1$  and  $f_2$  are unimodal,  $f_3$  and  $f_4$  are multimodal. Moreover,  $f_1$  and  $f_4$  are with separate variables,  $f_2$  and  $f_3$  are with linkage variables [3][4].

Table 1. Test functions for comparisons

Test function	Range	$f_{\min}$	Name
$f_1(x) = \sum_{i=1}^{D} x_i^2$	[-100,100] <sup>30</sup>	0	Sphere
$f_2(x) = \sum_{i=1}^{D} (\sum_{j=1}^{i} x_j)^2$	$[-100, 100]^{30}$	0	Quadric
$f_3(x) = \sum_{i=1}^{D-1} [100(x_{i+1} - x_i^2)^2 + (x_i - 1)^2]$	[-10,10] <sup>30</sup>	0	Rosenbrock
$f_4(x) = \sum_{i=1}^{D} -x_i \sin(\sqrt{x_i}) + 4189829 \times D$	$[-500, 500]^{30}$	0	Schwefel

Four DE variants, denoted as DE/best/0.1, DE/best/0.9, DE/rand/0.1, and DE/rand/0.9 are adopted for comparisons. Here DE/best and DE/rand indicate using the best mutation scheme and the random mutation scheme, while 0.1 and 0.9 indicate that CR=0.1 and CR=0.9. For CEDE, we set M=4 and the above four DE variants are adopted by each population respectively. Therefore, CEDE uses both the greedy and random mutation schemes, and also uses both a small CR value and a large CR value. All the DEs are with F=0.5 and the population size is set to 50. In CEDE, each population is with size of 50 and is dynamically controlled by AMS during the evolutionary process.

The maximal function evaluations (FEs) are  $3.0 \times 10^5$  for each D=30 dimensions function. Each function is simulated 30 times independently and their mean results are used in the comparison. Moreover, Wilcoxon's rank sum tests with significant level  $\alpha=0.05$  are conducted to make the results statistically sound. The experimental results are presented in Table 2.

Table 2	2. Results	compariso	ns with o	conventional	DEs
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Fu	nction	CEDE	DE/best/0.1	DE/best/0.9	DE/rand/0.1	DE/rand/0.9
$f_1$	mean	4.27E-139	3.24E-142-	2.93E-292-	3.26E-089+	9.04E-106+
	std	1.71E-138	8.94E-142	0.00E+000	3.25E-089	3.65E-105
$f_2$ I	mean	8.03E-027	1.15E+002+	1.67E+002+	2.33E+003+	9.28E-015+
	std	2.00E-026	4.98E+001	9.13E+002	4.79E+002	1.49E-014
$f_3 = \frac{\text{me}}{\text{st}}$	mean	5.96E-022	3.04E+001+	1.20E+000+	2.36E+001+	1.72E+001+
	std	3.25E-021	1.97E+001	1.86E+000	1.59E+000	6.11E+000
$f_4$	mean	3.82E-004	5.50E+002+	4.40E+003+	3.82E-004=	2.50E+003+
	std	0.00E+000	2.28E+002	7.47E+002	0.00E+000	6.03E+002
$(\pm)$ $(\pm)$ and $(\cdot)$ mean that CEDE performs significantly better than similar to and						

'+', '=', and '-' mean that CEDE performs significantly better than, similar to, and significantly worse than the corresponding DE, respectively.

The comparisons show that the performance of conventional DEs on different problems is seriously affected by the operators and/or parameters. For example, DEs use greedy mutation schemes yields much better results than DE use random mutation schemes on unimodal functions. However, when solving multimodal functions, greedy mutation seems to be inefficient, e.g., DE/best/0.1 and DE/best/0.9 are totally trapped by the multimodal functions  $f_3$  and  $f_4$ . Moreover, although using the same DE/best mutation scheme on  $f_1$ , DE/best/0.9 is remarkably better than DE/best/0.1. DE/rand/0.1 can obtain global optimum on  $f_4$  but DE/rand/0.9 is totally trapped. Therefore, it seems no a single DE mutation scheme and/or a single parameter setting can perform well on different kinds of problems. As CEDE uses multiple populations with different operators and different parameters, it is expected that CEDE is suitable for different kinds of problems. The results show that CEDE can obtain the global optimum of all 4 functions in all the 30 independent runs, no matter on unimodal or multimodal functions, no matter on functions with variable linkage or with variable independent.



process when optimizing the function.

Fig. 1 plots the dynamic population size during the search process. Fig. 1(a) shows that the population size of DE/best/0.9 keeps increasing during the evolutionary process on  $f_1$ . This is consistent with the fact that DE/best mutation scheme and large CR value have fast convergence speed to optimize simple unimodal function. However, when optimizing multimodal function, e.g.,  $f_4$ , as in Fig. 1(b), DE/best/0.9 performs very poor and therefore its population size keeps decreasing. Contrarily, the population sizes of DE/rand/0.1 and DE/best/0.1 increase during the evolutionary process. This is because that DE with small CR value can do well on separate multimodal function. Therefore, the figures show that AMS can catch the performance of different populations and let the individuals migrate to well-performed populations. This observation demonstrates the proposed AMS works well. Further study may test some other migration schemes as proposed by Zhong et al. [5] to evaluate the algorithm performance.

## 4. CONCLUSIONS

A CEDE with AMS is proposed. CEDE uses multiple populations to optimize the problem cooperatively, with each population using different operators and/or different parameters. When solving different kinds of problems, different populations may have different performance. Therefore, CEDE has promising performance on different problems. The experimental studies demonstrate this intuition and the good performance of CEDE.

## 5. ACKNOWLEDGMENTS

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