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A Hybrid Differential Evolution Algorithm for Solving Function Optimization

Online: 2010-06-07

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Keywords: Differential evolution (DE); evolutionary algorithm; optimization.

Abstract. One of the key points resulting in the success of differential evolution (Danis its mechanism of different mutation strategies for generating mutant vectors. In the paper, we also present a novel mutation strategy inspired by the velocity updating scheme of criticle system optimization (PSO). The proposed approach is called HDE, which conducts he mutatic strategy on the global best vector for each generation. Experimental studies on well-bown techmark functions show that HDE outperforms other three compared DE algorithms. Prost test cases.

Introduction

Differential Evolution (DE) [1] is a recently proposed of lutionary schpique, which has been shown to be a simple yet efficient evolutionary algorithm for many optimization problems [2]. Since the DE algorithm is simple, easy implementation are robustness it has been applied to many real-world problems, such as image processing, signal design.

However, the performance of DE is still quit to endent on the setting of control parameters such as the mutation factor and the crossover probability and ding to both experimental studies and theoretical analyses. Although there are some segestions for parameter settings, the interaction between the parameter setting and the optimization performance is still complicated and not completely understood. This is may be ause there is no fixed parameter setting that is suitable for different kinds of problems.

To tackle this problem rs proposed different parameter setting mechanisms. The any researc control parameter is altered some deterministic rules without taking into account any feedback from the evolutionar search. example is the time dependent change of the mutation rates proposed by Holland [3]. Gampen, et al. [4] evaluated different parameter settings for DE on the Sphere, Rosenbook's and Rastrigin's functions. Their experimental results revealed that the global optimum searching pability and the convergence speed are very sensitive to the choice of control Fathermore, a plausible choice of the population size ps is between 3D and F, and parameter 8D, with the saling fact, F = 0.6 and the crossover rate CR in [0.3, 0.9]. Recently, the authors in [5] F < 0.95 with F = 0.9 is a good first choice. CR typically lies in (0, 0.2) when claim th the function's separable, while in (0.9, 1) when the function's parameters are dependent. Liu and Lampinen [6] oposed a fuzzy adaptive DE (FADE), which uses a fuzzy logic controller to set the probability of mutation and crossover. Qin and Suganthan [7] presented a self-adaptive DE (SaDE) for numerical optimization, which focused on adaptation for parameter CR and mutation strategies of DE. Brest et al. [8] introduced self-adapting control parameter settings in DE (SADE). Yang et al. [9] introduced a neighborhood search strategy to DE (NSDE), which generates F from Gaussian and Cauchy distributed random numbers instead of predefining a constant F. Ali and Torn [10] introduced auxiliary population and automatic calculating of the amplification factor F.

In this paper, we propose a hybrid DE algorithm to improve the performance of DE. The proposed approach is called HDE, which employs a novel mutation strategy inspired by the velocity updating model of PSO. Experimental studies on eight well-known benchmark problems show that HDE obtains better performance when comparing with other three DE variants.

Differential Evolution

The main idea of DE is not to mutate vector components by simply replacing their values by random values. Instead, two population mates are randomly selected whose weighted difference is added to a third randomly selected population member creating this way a mutant vector. Then, either a two-point crossover [1] or a multi-point crossover [1] is performed between the mutant vector and the current population member being considered. At the end, the new created offspring vector (trial vector) replaces the current considered vector in the next generation, if its fitness is better. Otherwise, the trial vector is discarded.

Let *D* and *ps* be the dimension of the problem and the population size, respectively. A vector member in the population can be defined by

$$X_{i,G} = (X_{1i,G}, X_{2i,G}, \dots, X_{Di,G})$$
(1)

where i = 1, 2, ..., ps, $G = 1, 2, ..., MAX_G$ and MAX_G is the maximum number of generations.

Like genetic algorithms, DE also has three operators, mutation, crossover and relection first, we create a mutant vector $V_{i,G}$. And then we recombine the mutant vector and a covernt vector to create a new trail vector $U_{i,G}$. At last, we compare the fitness value of $V_{i,G}$ with $C_{i,G}$, an select obetter one as the new current vector in the next generation. The three operators and describes a follows.

DE uses the difference between randomly selected individuals as a resource of random variations for a third individual, referred to as the target individual f is solution are generated by adding weighted differences vector. This process is referred as a mutation operator described as follows. For each vector $X_{i,G}$ in generation G, a mutant vector $V_{i,G}$ is refined by

$$V_{i,G} = X_{r1,G} + F(X_{r2,G} - X_{r3,G}) \tag{2}$$

where $X_{i,G}$ (i = 1, 2, ..., ps) are solution vectors in general G, ps is the population size, i = 1, 2, ..., ps and r_1 , r_2 , and r_3 are mutually different random integer in lines selected from $\{1, 2, ..., ps\}$.

After the mutation step, a new training $U_{i,G} = V_{1i,G}, U_{2i,G}, ..., U_{Di,G}$ is generated by recombination of the mutant vector and the current ectors.

$$U_{ji,G} = \begin{cases} V_{ji,G}, & \text{if } rand \text{ if } v \leq CR \lor j = k \\ X_{ji,G}, & \text{other ise} \end{cases}$$
(3)

where CR is the predefined cross per probability, and $rand_{j}(0,1)$ is a random number within (0, 1) for the *i*th dimension and $\{0, 1, 2, ..., D\}$ is a random parameter index.

After the two stars a selection mechanism is used to choose a better vector between $U_{i,G}$ and $X_{i,G}$ to update a current ector in the next generation. For a minimization problem, the vector with smaller fitness value is better.

The Proposition Approach

Generally, the termination condition of an evolutionary algorithm is that the global best individual in the population finds the global optimum or the number of function evaluations reaches to the predefined maximum value. It can be found that the global best individual is important toward searching the global optimum. If the global best individual is trapped, the whole population may not find good solutions. To tackle this problem, we propose a novel mutation strategy on the global best individual. The motivation of the approach is inspired by the velocity updating model of PSO which utilizes the searching experiences of the previous best individuals and the global best individual.

In PSO, an individual in the population is called "particle". Each particle has two vectors: position and velocity, which are updated as follows [11].

$$V_{i}(t+1) = w * V_{i}(t) + c_{1} * rand1() * (pbest_{i} - X_{i}(t)) + c_{2} * rand2() * (Best - X_{i}(t))$$

$$\tag{4}$$

$$X_{i}(t+1) = X_{i}(t) + V_{i}(t+1)$$
(5)

where X_i and V_i are the position and velocity of the *i*th particle, *pbest_i* and *Best* are previous best particle of the *i*th particle and the global best particle found by all particles so far respectively, w is an inertia factor, rand1() and rand2() are two random numbers independently generated within the range of [0,1], and c_1 and c_2 are two learning factors.

Based on the velocity updating equation, we propose a novel mutation strategy as follows.

$$Best^* = a_1 * Best + a_2 * (Best - X_{i1}) + a_3 * (X_{i2} - X_{i1})$$
(6)

where Best is the global best individual in the population, X_{i1} and X_{i2} are two different random number within [1, ps], $i1 \neq i2$, ps is the population $[1, a_2, a_3]$ are three random numbers within [0,1], and $a_1 + a_2 + a_3 = 1$. The random numbers can generate by

$$a_1 = rand(0,1), a_2 = rand(0,1), a_3 = rand(0,1)$$
 (7)

$$sum = a_1 + a_2 + a_3 \tag{8}$$

$$a_1 = a_1/sum, a_2 = a_2/sum, a_3 = a_3/sum$$
 (9)

where rand(0,1) is a random number within [0,1].

The Framework of HDE orithmeter

```
while NE < MAXNE
    for i = 1
                es do
                  trial indevidual according to Eq. 2 and Eq. 3;
       Ge derate
                 the fitness of the trial individual;
               fitter one between Xi and the trail;
        Jelec
     end for
        perate random numbers according to Eq. 7, Eq. 8 and Eq. 9;
         ct the mutation according to Eq. 6;
    If Be is better than Best
       Replace Best with Best*;
     nd if
  end while
End
```

Every receive we conduct the mutation strategy. If the individual $Best^*$ after the mutation is better than toold Best, then replace the Best with $Best^*$; otherwise keep the Best unchangeable. The application of the mutation strategy used in DE is described in the framework of the HDE algorithm, where ps is the population size, Best is the global best individual in the population, NE is the number of function evaluations, and MAX_{NE} is the maximum number of function evaluations.

Simulation Studies

Test Functions. In this paper, we test the proposed approach HDE on eight well-known benchmark problems, which were used in early studies [12]. All the functions used in this paper are to be minimized. The description of the benchmark functions and their global optima are listed as follows.

$$f_1 = \sum_{i=1}^D x_i^2$$

where $x_i \in [-100, 100]$, D=30, and the global optimum is 0.

$$f_2 = \sum_{i=1}^{D} |x_i| + \prod_{i=1}^{D} x_i$$

where $x_i \in [-10, 10]$, D=30, and the global optimum is 0.

$$f_3 = \sum_{i=1}^{D} \left(\sum_{j=1}^{i} x_j\right)^2$$

where $x_i \in [-100, 100]$, D=30, and the global optimum is 0.

$$f_4 = \max_{i} (|x_i|, 1 \le i \le D)$$

where $x_i \in [-100, 100]$, D=30, and the global optimum is 0.

$$f_5 = \sum_{i=1}^{D-1} \left[100(x_{i+1} - x_i^2)^2 + (x_i - 1)^2 \right]$$

where $x_i \in [-30, 30]$, D=30, and the global optimum is 0.

$$f_6 = \sum_{i=1}^{D} \left(\left\lfloor x_i + 0.5 \right\rfloor \right)^2$$

where $x_i \in [-100, 100]$, D=30, and the global optimum is 0.

$$f_7 = \sum_{i=1}^{D} ix_i^4 + rand[0,1)$$

where $x_i \in [-1.28, 1.28]$, D=30, and the global optimum is 0

$$f_8 = \sum_{i=1}^{D} -x_i \sin\left(\sqrt{|x_i|}\right)$$

where $x_i \in [-500, 500]$, D=30, and the global optimum is

Comparison of HDE with DE. In this section, we compare the proposed approach HDE with classical DE on the ten test problems. In the sake of fair competition, we use the same parameter settings for HDE and DE. The parameter size ps, QR and P are set to 50, 0.9 and 0.5, respectively. Both DE and HDE use the same multiple strengly QE/rand/1. All the experiments in this paper are conducted 30 times with different random seeds, and the average results throughout the optimization runs are recorded.

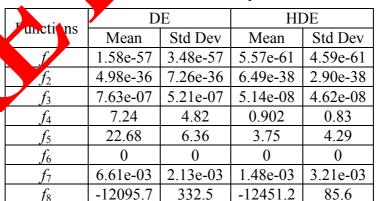


Table The results achieved by DE and HDE.

The comparison results between HDE and DE are presented in Table 1, where "Mean" indicates the mean best function values, and "Std Dev" stands for the standard deviation. From the results, it can be seen that HDE outperforms DE in all test cases except for functions f_6 . On this function, both HDE and DE achieve the same results.

Comparison of HDE with SaDE and NSDE. This section presents another comparative study to further verify the performance of HDE. The involved algorithm includes Self-adaptive DE (SaDE)



and DE with neighborhood search (NSDE). The parameter settings of SaDE and NSDE are described in [13]. For IDE, the population size ps, CR and F are set to 50, 0.9 and 0.5, respectively. The initial population is generated uniformly at random in the search domain of the functions. The maximum number function evaluations (MAX_{NE}) are listed in Table 2.

Functions	MAX _{NE}	SaDE	NSDE	HDE
		Mean	Mean	Mean
f_1	150000	7.49e-20	7.76e-16	5.57e-61
f_2	150000	6.22e-11	4.51e-10	6.49e-38
f_3	150000	1.12e-18	1.06e-14	5.14e-08
f_4	150000	2.96e-02	2.54e-02	0.902
f_5	500000	2.10e+01	1.24e+01	3.75
f_6	150000	0	0	0
f_7	150000	7.58e-03	1.20e-02	1.4 se-03
f_8	150000	-12569.5	-12569.5	1. 11.2

Table 2. The results achieved by SaDE, NSDE and HDE.

The average results over 30 trails of HDE, SaDE and NSDE are presented in Trole 2, where "Mean" indicates the mean best function values, and "Std Dev" state for the state of deviation. The results of SaDE and NSDE are taken form Table 2 and 3 in [13] Front the results at cane be seen that HDE outperforms SaDE and NSDE on functions f_1 , f_2 , f_5 and f_5 and sight vantly improve the results on functions f_2 and f_5 . For function f_6 , all the three algorithms obtain the tame performance. SaDE achieves better results than NSDE and HDE on function f_3 , while NSDE outperforms the other two DE algorithms on function f_4 . Both SaDE and NSDE across better results than IDE on function f_8 .

Conclusion

This paper presents a hybrid DE algorithm to improve the performance of DE. The proposed approach is called HDE, which employs a novertestation strategy inspired by the velocity updating model of PSO. Every generation, we conduct the metation on the global best individual, and select a better one between the mutant and the best as recent best individual. Experimental studies on eight well-known benchmark problems show that HDE obtains better performance than classical DE, SaDE and NSDE in most decrease.

However, HDE and other toppared DE algorithms falls into local minima on function f_5 . It suggests that the proposed mutative strategy is not suitable for all kinds of problems. It is worth to introduce more officient trategies into DE to improve its performance in the future work.

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