

# An Efficient Constraint Handling Approach for Optimization Problems with Limited Feasibility and Computationally Expensive Constraint Evaluations

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## ABSTRACT

Existing optimization approaches adopt a *full evaluation policy*, i.e. all the constraints corresponding to a solution are evaluated throughout the course of search. Furthermore, a common sequence of constraint evaluation is used for all the solutions. In this paper, we introduce a scheme of constraint handling, wherein every solution is assigned a random sequence of constraints and the evaluation process is aborted whenever a constraint is violated. The solutions are sorted based on two measures i.e. the number of satisfied constraints and the violation measure. The number of satisfied constraints takes a precedence over the amount of violation. We illustrate the performance of the proposed scheme and compare it with other state-of-the-art constraint handling methods within a framework of differential evolution. The results are compared using *g-series* test functions for inequality constraints. The results clearly highlight the potential savings offered by the proposed method.

## Categories and Subject Descriptors

I [Computing Methodologies]: MISCELLANEOUS

## General Terms

Algorithm

## Keywords

Constraint Handling, Constraint Sequencing, Fitness Evaluation

## 1. INTRODUCTION

Constraint handling is an important area of research and various forms of constraint handling schemes have been proposed in literature. The performance of all population-based stochastic optimization algorithms are known to be affected by the presence of constraints. The nonlinearity, multi-modality and the feasibility space associated with each constraint is likely to be different. Constraint handling methods can be broadly categorized in four different types i.e. use of penalty functions, repair schemes, use of decoders and the separation of objective function and constraints [1]. More recent methods maintain infeasible solutions such as through stochastic ranking,  $\epsilon$  based comparisons or adaptive penalty function formulations [2]. However, in all such formulations and implementations, a *full evaluation policy* is adopted, i.e. for every solution, its constraint violation (CV) measure is computed which is the sum

of all constraint violations. An important question is “why do we spend computational resources to evaluate constraints of a solution, when it has already violated a constraint?”. Assuming that one is only interested in a feasible solution (preferably optimum) at the end of the search process, it is important to investigate the worth of evaluating infeasible solutions i.e. the cost of evaluation versus the knowledge gained to steer the search. Other followup questions include “what is the best sequence to evaluate the constraints?” and “is there a benefit in using different sequence of constraints?”. This paper attempts to understand the cost-benefit of *partial evaluation policy* i.e. aborting evaluation of constraints if the solution has already violated one constraint. Above discussion becomes more relevant in the context of optimization problems involving computationally expensive constraint evaluations. The study assumes that the constraints can be evaluated independently of one another.

An optimization algorithm has been introduced based on a partial evaluation policy. The solutions in the population are evaluated based on a random sequence of constraints. The search using multiple constraint sequences offer the potential to reach different regions of the search space. The proposed scheme has been implemented using a framework based on differential evolution [3].

## 2. PROPOSED ALGORITHM

A population of  $N$  individuals is initialized. The variables of  $i^{\text{th}}$  individual are initialized as follows:

$$x_{j,i} = x_{j,\min} + \text{rand}_{i,j}[0,1) \cdot (x_{j,\max} - x_{j,\min}) \quad (1)$$

where  $j = 1, 2, \dots, D$  is the number of variables;  $x_{j,\max}$  and  $x_{j,\min}$  are the upper and the lower bounds of  $j^{\text{th}}$  variable. For a problem with  $m$  constraints, each individual is assigned a random sequence of constraints for evaluation. Every individual of the population is evaluated using its prescribed constraint sequence. Whenever a constraint is violated, the evaluation is aborted. The term *number of function evaluations* referred in the paper is the sum of the number of evaluated constraints and objective function evaluations [4].

In order to generate an offspring solution, the first parent is selected sequentially, the second and third parents are selected randomly from the entire population. In the recombination process, a binomial crossover [2] has been used to generate the offspring solution. The fitness of a solution is determined as follows:

$$\text{fitness}(\xi) = \begin{cases} f(\mathbf{x}), & \mathbf{x} \in \mathbb{R}^n \\ c_i, & i = 1, 2, \dots, m \end{cases} \quad (2)$$

where  $c_i$  is the constraint violation measure of  $m$  number of constraints. The equality constraints are transformed into a set of inequalities as  $|h_j(\mathbf{x}) - \delta| \leq 0$  (assuming  $\delta$  is small positive quan-

**Algorithm 1** DE-CS

**SET:**  $NT_{max}$ {Total number of function evaluation},  $N$ {Size of population},  $CR$ {Crossover rate},  $F$ {A Mutation scale factor},  $Evalcount = 0$

- 1: Initialize the population of individuals and assign a random constraint sequence to each individual;
- 2: Evaluate the solutions following the above assigned sequence of constraints;  $Update(Evalcount)$ ;
- 3: **while** ( $Evalcount \leq NT_{max}$ ) **do**
- 4:   **for**  $i=1:N$  **do**
- 5:     Select  $P_1 = i$  i.e. the  $i^{th}$  parent and two other parents  $P_2$  and  $P_3$  randomly s.t.  $P_1 \neq P_2 \neq P_3$ ;
- 6:     Generate an offspring using recombination;
- 7:     Evaluate the offspring using the sequence of  $P_1$ ;  $Update(Evalcount)$ ;
- 8:     The offspring is compared with solutions in the population for replacement based on fitness
- 9:   **end for**
- 10: **end while**

\* $Evalcount$  denotes the sum of objective and all individual constraint evaluations

tity). Here,  $c_i$  denotes the constraint satisfaction vector where

$$c_i = \begin{cases} 0, & \text{if } i^{th} \text{ constraint satisfied } i = 1, 2, \dots, m \\ g_i(\mathbf{x}), & \text{if } i^{th} \text{ constraint violated, } i = 1, 2, \dots, q \\ |h_j(\mathbf{x}) - \delta|, & \text{if } i^{th} \text{ constraint violated, } i = q + 1, \dots, m \end{cases} \quad (3)$$

For every solution in the population, one can compute the number of satisfied constraints ( $NS$ ) and the amount of violation ( $V$ ). With the number of constraints satisfied taking a precedence over the violation value, a sorting would yield the ranks of the individual solutions. For example assume a population, containing 4 solutions ( $S1, S2, S3, S4$ ). The constraint violation matrix would assume a form illustrated in Table 1 with  $S3$  identified as the best and  $S1$  the worst.

**Table 1: Ranking of 4 individuals in the population in presence of 3 constraints**

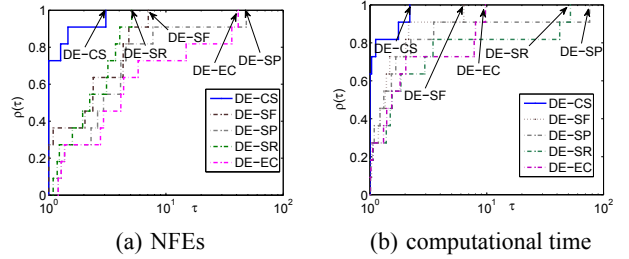
Initial order				$NS$	$V$	Final rank
$S1(g_1, g_2, g_3)$	5	—	—	0	5	4
$S2(g_2, g_3, g_1)$	0	3	—	1	3	2
$S3(g_1, g_3, g_2)$	0	0	1	2	1	1
$S4(g_2, g_1, g_3)$	2	—	—	0	2	3

### 3. EXPERIMENTAL RESULTS

The above section illustrated the principles of constraint sequencing and partial evaluation. In this section we objectively evaluate its performance on *CEC-2006* [5] benchmarks. We also include the results obtained by using stochastic ranking (SR) [6], self adaptive penalty (SP) [7], superiority of feasibility (SF) [8] and epsilon constraint (EC) [9] within the same framework of DE. Results based on performance profile are included for a more objective comparison. A population size of 50 is used for all the problems and the results are computed based on 30 independent runs. A fixed value of  $CR = 0.9$  and  $F = 0.5$  have been set for all the cases resulting the number of function evolutions (i.e.  $NFEs$ ) of  $4800 * (N * m)$ , where  $N$  is the size of the population and  $m$  is the number of constraints.

In this experiment, we observe how quickly a feasible solution appears in the population—the function evaluation to reach a feasible solution and the computational time required to achieve the feasible solution. A performance profile [10] is computed for a more objective comparison between the strategies. The results clearly indicate the superiority of DE-CS over other strategies in terms of

$NFEs$  and computational time. Figure 1 shows the value of  $\rho(\tau)$  for  $r_{p,s} \leq \tau$  of the normalized performance ratio [10] i.e. (a) the number of function evaluation (b) computational time. One can observe from the figure that DE-CS outperforms with other strategies in terms of both.



**Figure 1: Performance profile of DE-CS and others**

### 4. CONCLUSION

In this paper, a scheme of constraint handling has been introduced within the framework of differential evolution utilizing the concepts of partial evaluation and constraint sequencing. The performance of the algorithm is subsequently assessed on 11 well known constrained single objective optimization benchmarks. The results on the test problems clearly indicate that the approach is computationally efficient and better than existing strategies for constraint handling.

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