

# SELF-ADAPTATION OF PARAMETERS FOR MCPC IN GENETIC ALGORITHMS

ESQUIVEL S.C., LEIVA H. A., GALLARD, R.H.

Proyecto UNSL-338403<sup>1</sup>  
Departamento de Informática  
Universidad Nacional de San Luis (UNSL)  
Ejército de los Andes 950 - Local 106  
5700 - San Luis, Argentina.  
E-mail: {esquivel, aleiva, rgallard}@unsl.edu.ar  
Phone: + 54 652 20823  
Fax : +54 652 30224

## Abstract

As a new promising crossover method, multiple crossovers per couple (MCPC) deserves special attention in evolutionary computing field. Allowing multiple crossovers per couple on a selected pair of parents provided an extra benefit in processing time and similar quality of solutions when contrasted against the conventional single crossover per couple approach (SCPC). These results, were confirmed when optimising classic testing functions and harder (non-linear, non-separable) functions.

Despite these benefits, due to a reinforcement of selective pressure, MCPC showed in some cases an undesirable premature convergence effect. In order to face this problem, the present paper attempts to control the number of crossovers, and offspring, allowed to the mating pair in a self-adaptive manner.

Self-adaptation of parameters is a central feature of evolutionary strategies, another class of evolutionary algorithms, which simultaneously apply evolutionary principles on the search space of object variables and on strategy parameters. In other words, parameter values are also submitted to the evolutionary process. This approach can be also applied to genetic algorithms.

In the case of MCPC, the number of crossovers allowed to a selected couple is a key parameter and consequently self-adaptation is achieved by adding to the chromosome structure “labels” describing the number of crossover allowed to each individual. Labels, which are bit strings, also undergo crossover and mutation and consequently evolve together with the individual. During the stages of the evolution process, it is expected that the algorithm will return the number of crossovers for which the current population exhibits a better behaviour.

Descriptions of different self-adaptation methods used, experiments and some of the results obtained are shown.

**Keywords:** genetic algorithms, self-adaptation, crossover, function optimisation.

---

<sup>1</sup> The Research Group is supported by the Universidad Nacional de San Luis and the ANPCYT (National Agency to Promote Science and Technology).

## SELF-ADAPTATION OF PARAMETERS FOR MCPC IN GENETIC ALGORITHMS

### 1. INTRODUCTION

MCPC as proposed in [1] allows multiple children per couple by replicated application of crossover. In those experiments the number of crossover allowed to a couple remain fixed during a single run. The rationale behind this implementation was to isolate MCPC effects to obtain a set of preliminary results. Previous experiments showed a quality of results as good as under SCPC, and sometimes better, when 3 and 4 crossovers per couple were allowed. Also an extra benefit in processing time was detected. Despite these benefits, due to a reinforcement of selective pressure, MCPC showed in some cases an undesirable premature convergence effect.

Self-adaptation is a new field in evolutionary computation which advises to dynamically update parameters of the algorithm by evolving them as part of the chromosome structure. Previous work of Spears [2] suggested adaptive approaches to select the type of crossover operator to be applied to each couple during a genetic algorithm execution. In this paper we propose a self-adaptive approach to determine the number of crossovers to be applied to a selected couple under MCPC. A general classification scheme, the criteria to implement self-adaptation when optimizing hard testing functions a description of experiments and results are shown in the following sections.

### 2. PARAMETER CONTROL: A CLASSIFICATION SCHEME

Today a great interest exists in methods including mechanisms to control parameters used by evolutionary algorithms during execution. Eiben, Hinterding and Michalewicz [3] gave the following main categories of parameter control:

- *Deterministic Parameter Control*: This is the case in which the parameter value is modified according with a deterministic rule, without any feedback of the searching process performed by the strategy.
- *Adaptive Parameter Control*: In this case some feedback information of the searching process is used to determine the direction and magnitude of the change in the parameters.
- *Self-adaptive Parameter Control*: Here the parameters to be adapted are codified within the chromosomes and undergo genetic operations. The best individuals of the population have better chances of survival and reproduction. Hence it is expected that better parameter values be more intensively propagated.

As the number of crossovers to be applied, to a couple, in MCPC is one of the parameters of the algorithm that is included as a part of an individual, our present approach appertains to the last above-mentioned category.

### 3. SELF-ADAPTIVE PARAMETER CONTROL IMPLEMENTATION

As we previously said, we attempt to self-adapt the number of crossovers per couple in MCPC. Because we are using a binary representation of chromosomes, the number of crossovers allowed for an individual is codified in a field at the rightmost positions of the bit string. Let us call it the *ncross\_field*. In some experiments we allowed a maximum of three and in others a maximum of seven crossovers per couple. So, two or three extra bits were enough for that purpose. More generally the last  $\log_2(max\_cross + 1)$  bits of each individual are used to find an expected optimum number of crossovers.

In that way we have two searching spaces: one corresponding to the objective function and other associated to the number of crossovers to apply.

Our attempt is that the individuals preserve the information about the number of crossovers originally applied to their parents. In this way it is expected that, based on the *survival-of-the-*

*fittest* principle, good solutions carry information about the number of crossover applied to their ancestors and that this number would be an appropriate one.

According to Spears [2] we used a local adaptive technique. Once the couple was selected we check the corresponding number of crossover carried by each parent and;

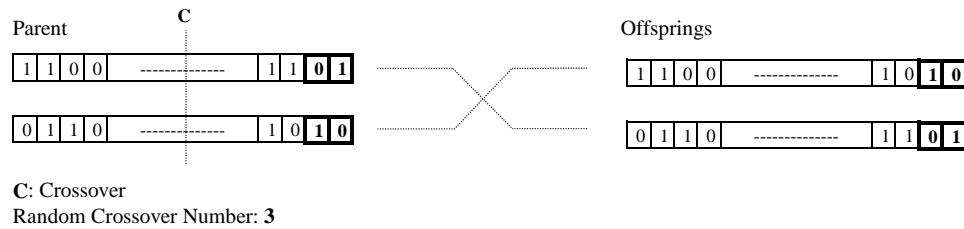
- If they match, then we apply the recombination operator a number of times specified by the *ncross\_field*.
- Otherwise we choose a random number in the permitted range.

In the second situation and following the Spears' approach, when decoded numbers of crossovers are different, we are violating our attempt to preserve information because the children do not keep the number of crossover by which they were created. If the crossover point does not disrupt the *ncross\_field* (and this event has low probability of occurrence) then children retain information from either parent, but they do not preserve information about how they were created.

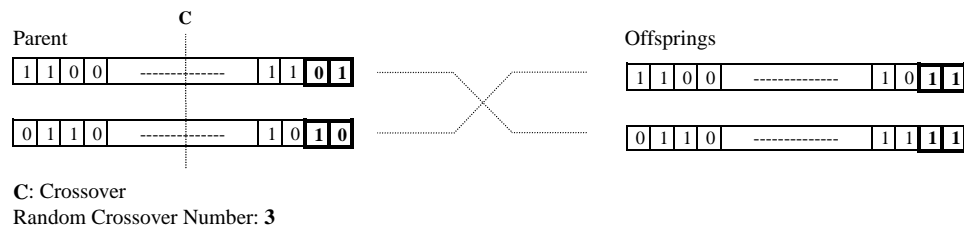
In order to either retain information about how an individual was created or how their parents were created, we devised two different approaches for experimentation.

- *E1*: In any situation, exchange of information from parents to children is done in the traditional way applying the genetic operators with their corresponding probabilities. Parent's chromosomes are mated and undergo crossover a certain number of times according to the specified values in *ncross\_field* if they match, or to a random allowed value otherwise. After recombination, mutation is applied to the children.

In the *don't match* situation, this approach, preserving parent's information, enforces population diversity in the parameter searching space, because most of the time one child inherits characteristics from one of the parent and the other child inherits features from the other parent.



- *E2*: If the values specified in *ncross\_field* do not match then the new random value for the number of crossovers is inserted first in the parent's *ncross\_field*, and afterwards crossover is performed for the number of times specified by this random value. This approach by preserving individual information creates more similar individuals in the parameter searching space and increases loss of genetic diversity.



#### 4. EXPERIMENTAL TESTS

Experiments E1 and E2, to verify the parameter control mechanisms were designed. For this we chose two hard testing functions: Easom's [5] and Schwefel's [6] functions (see table 1). A modified version of the non-canonical genetic algorithm (Goldberg [4]) was implemented.

Series of many runs were performed on each function, with randomised initial population of size fixed to 1000 individuals, using binary representation, proportional selection, elitism, one point crossover and bit flip mutation. The number of generations was fixed to 500 and probabilities for crossover and mutation were fixed to 0.65 and 0.05, respectively.

The relatively big population size of 1000 individuals was chosen in order to allow a significant contribution of selected individuals to the evolution process when high number of crossovers are allowed. For example in the case of six crossovers per couple only 16.5% of the available individuals in the old population will intervene when building the new generation.

Notation	Description	Characteristics
<i>f5: Easom's Function</i>	$f_5(x_1, x_2) = -\cos(x_1) \cos(x_2) e^{-((x_1-\pi)^2 + (x_2-\pi)^2)}$ , $x_1, x_2 \in [-100, 100]$	Unimodal, the global minimum has a small area relative to the search space
<i>f7: Schwefel's Function 7</i>	$f_7(\vec{x}) = \sum_i -x_i \cdot \sin(\sqrt{ x_i })$ , for $i = 1 : n$ $x_i \in [-500, 500]$	Dim. $N = 5$ Highly multimodal, the global minimum is geometrically distant from the next best local minima.

Table 1. Objective functions

The following variables were chosen for the analysis

Name	Description
<b>Quality</b>	Is the ratio $\left(\frac{best\_value}{opt\_value}\right)$ between the best value and the optimal value. It gives a measure for the quality of a solution.
<b>CrAvg</b>	Mean number of crossover allowed per selected couple.
<b>Dtime</b>	Defined as in [1]. Running time difference. It is the percentile of time reduction when compared with classic crossover (single crossover per couple).

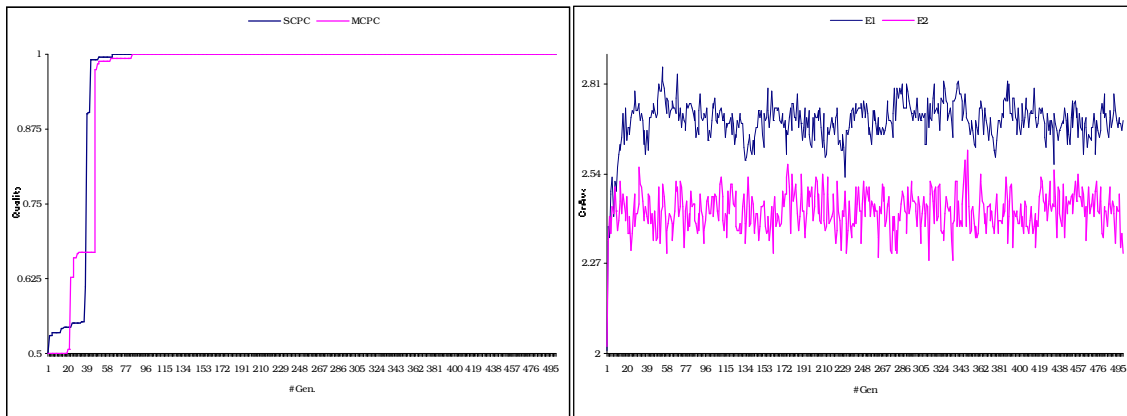
#### 5. RESULTS

Results concerning *Quality* were similar under E1 and E2, but different when *CrAvg* is considered. Hence the following figures alternatively show *Quality* results under experiments E1 or E2. Results in regard to *CrAvg* are summarised in the same graph for both experiments.

##### *Function f5*

We started optimizing *f5*, and for this unimodal function two bits were used to code the *ncross\_field*, allowing a maximum of three crossovers (six children per couple).

In figure 1 *Quality* values for the Easom's function, show a slightly slower convergence of MCPC when compared with SCPC, but after 80 generations the former results are better. In fact during the simulation, values of *Quality* reach 1.0 under MCPC and 0.9998 under SCPC.



**Fig. 1: Quality and  $CrAvg$  values for function  $f_5$**

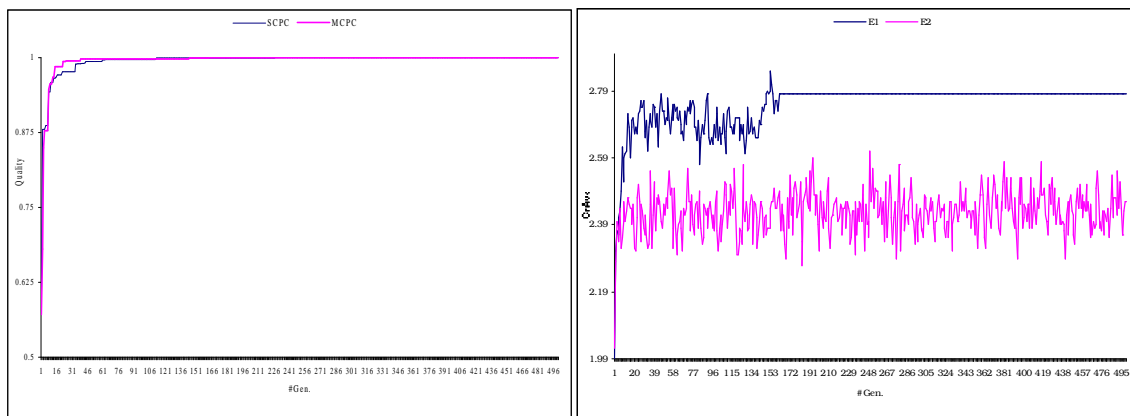
Regarding  $CrAvg$  values, it can be observed that they oscillate between 2.5 and 2.8 after the few first generations under E1 and between 2.3 and 2.6 under E2.

Here the behaviour of the self adaptive parameter control mechanism is clear: when genetic diversity in the parameter searching space is low then lesser number of crossovers are allowed and viceversa. This behaviour favours the evolutionary process.

Concerning to  $Dtime$  values they were 37,7% and 33,6% under E1 and E2 approaches respectively.

### Function $f_7$

When optimizing  $f_7$ , we decided to use two and also three bits to code the  $ncross\_field$ , allowing a maximum of three and seven crossovers respectively.



**Fig. 2: Quality and  $CrAvg$  values for function  $f_7$**

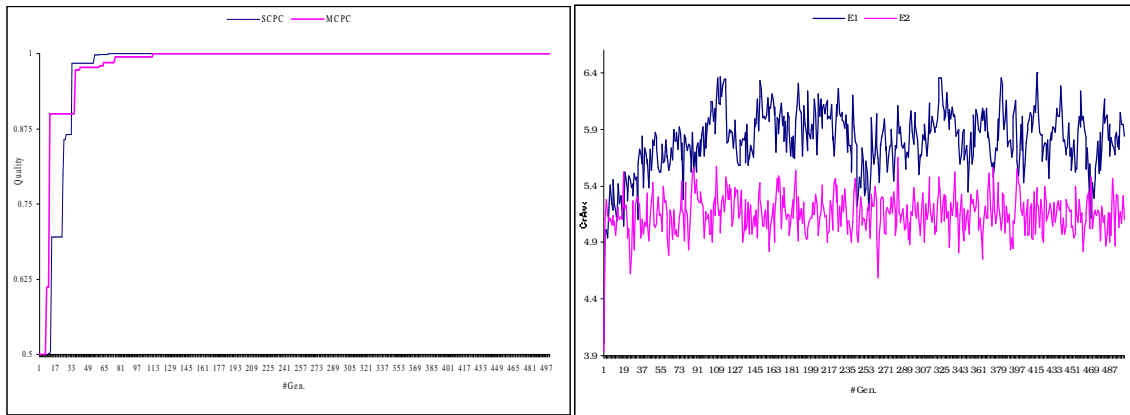
In figure 2  $Quality$  values for the Schwefel's function with two bits for  $ncross\_field$ , show a slightly faster convergence of MCPC when compared with SCPC, but both converge to the optimum reaching a  $Quality$  value of 1.0.

Regarding  $CrAvg$  values, it can be observed that they oscillate between 2.6 and 2.8 remaining stable after 152 generations under E1, and oscillate between 2.3 and 2.6 under E2.

Here again the behaviour of the self-adaptive parameter control mechanism is shown. Parameter setting is adapted to the population diversity in the parameter searching space. In this case this behaviour prevent the evolutionary process of being trapped in a local optima.

Concerning to  $Dtime$  values they were 22.1% and 21.3% under E1 and E2 approaches respectively.

Another set of experiments studied the behaviour of the control mechanism for the Schwefel's function with three bits for the *ncross\_field*.



**Fig. 3:** *Quality* and *CrAvg* values for function  $f_5$

In figure 3 *Quality* values for the Schwefel's function with three bits for *ncross\_field*, are shown. Here we cannot clear differences on convergence velocity. At the beginning MCPC shows to be faster and after that SCPC is faster 110 generations both reach the optimum.

Regarding *CrAvg* values, it can be observed that they oscillate between 4.5 and 5.6 under E1, and between 5.2 and 6.3 under E2.

Here again the behaviour of the self-adaptive parameter control mechanism is shown. Parameter setting is adapted not only to the population diversity but to the maximum number of crossovers allowed. Concerning to *Dtime* values they were 50% and 48% under E1 and E2 approaches respectively.

## 6. CONCLUSIONS

The present paper proposes an alternative approach to assign the number of crossovers allowed for a selected couple. Instead of doing that based on the couple fitness as in [8], here we suggest to use a self-adaptive parameter control approach. The parameter considered is coded in a *ncross\_field* of two or three bits in the chromosome structure, and further submitted to genetic operations in the same way as any evolutionary technique does. Within this approach two searching processes are carried out simultaneously: one on the problem (objective function) space and the other on the parameter space. In this way it is expected an adaptive parameter setting, retaining best settings through the survival of the fittest individuals in the problem space.

Two different strategies were approached in order to overcome loss of information about offspring or parents creation. As they were conceived, approach *E1* maintains population diversity in the parameter searching space while approach *E2* leads to a loss of diversity.

Being consequent with this situation the control mechanism adapts the number of crossovers for exploration (under *E1*) or exploitation (under *E2*) accordingly. And this behaviour is preserved for diverse maximum number of crossover allowed.

It is also remarkable that on each experiment MCPC outperforms SCPC most of the time on quite different fitness landscapes.

Future work will consider more biased methods, tied to the fitness of individuals in the couple, to choose the number of crossovers in the *don't-match* case.

## 7. ACKNOWLEDGEMENTS

We acknowledge the cooperation of the project group for providing new ideas and constructive criticisms. Also to the Universidad Nacional de San Luis the CONICET, and the ANPCYT from which we receive continuous support.

## 8. BIBLIOGRAPHY

- [1] Esquivel S., Leiva A., Gallard R.: *Multiple Crossover per Couple in Genetic Algorithms*. Proceedings of the Fourth IEEE Conference on Evolutionary Computation (ICEC'97), Indianapolis, USA, April 1997.
- [2] William M. Spears: *Adapting Crossover in Evolutionary Algorithms*. Proceedings of the Evolutionary Programming Conference, 1995.
- [3] Agoston E. Eiben, Robert Hinterding, Zbigniew Michalewicz: *Parameter Control in Evolutionary Algorithms*. Technical Report, UNC - Charlotte, 1998.
- [4] Goldberg D.E.: *Genetic Algorithms in Search Optimization and Machine Learning*. Addison Wesley Publishing, 1989.
- [5] Easom, E.: *A survey of global optimization techniques*. M. Eng. Thesis, Univ. Louisville, Louisville, KY, 1990. In [7].
- [6] Schwefel, H. P.: *Numerical optimization of computer models*. Chichester: Wiley & Sons, 1981. In [7].
- [7] Hartmut Pohlheim: *Genetic and Evolutionary Algorithm Toolbox for use with Matlab (GEATbx)*. Copyright © 1996, Germany. All rights reserved.
- [8] Esquivel S., Leiva A., Gallard R.: *Couple Fitness Based Selection with Multiple Crossover per Couple in Genetic Algorithms*. Proceedings of the International Symposium on Engineering of Intelligent Systems (EIS'98), La Laguna, Tenerife, Spain, February 1998.