

# Evolutionary Multiobjective Optimization using a Cultural Algorithm

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**Abstract**—In this paper, we present the first proposal to use a cultural algorithm to solve multiobjective optimization problems. Our proposal uses evolutionary programming, Pareto ranking and elitism (i.e., an external population). The approach proposed is validated using several examples taken from the specialized literature. Our results are compared with respect to the NSGA-II, which is an algorithm representative of the state-of-the-art in evolutionary multiobjective optimization. The performance of our approach indicates that cultural algorithms are a viable alternative for multiobjective optimization.

**Keywords**—cultural algorithms, multiobjective optimization, evolutionary programming.

## I. INTRODUCTION

In recent years, evolutionary algorithms have been widely used for multiobjective optimization, obtaining very promising results [3], [2]. An important advantage of evolutionary algorithms over traditional techniques used for multiobjective optimization is that the former operate over a set of solutions at a time. Therefore, evolutionary algorithms can produce several elements of the Pareto optimal set in a single run rather than generating one at a time as traditional mathematical programming approaches.

Cultural algorithms [18] are a technique that incorporates domain knowledge obtained during the evolutionary process as to make the search process more efficient. Cultural algorithms have been successfully applied to several types of optimization problems (e.g., in constrained single-objective optimization problems [20], [12], [13], [4]). However, until now, nobody had proposed a cultural algorithm for multiobjective optimization adopting Pareto ranking and elitism. Ours is then the first proposal in this direction.

## II. BASIC CONCEPTS

**Definition 1 (General MOP):** Find the vector  $\vec{x}^* = [x_1^*, x_2^*, \dots, x_n^*]^T$  which will satisfy the  $m$  inequality constraints:

$$g_i(\vec{x}) \geq 0 \quad i = 1, 2, \dots, m \quad (1)$$

the  $p$  equality constraints

$$h_i(\vec{x}) = 0 \quad i = 1, 2, \dots, p \quad (2)$$

and will optimize the vector function

$$\vec{f}(\vec{x}) = [f_1(\vec{x}), f_2(\vec{x}), \dots, f_k(\vec{x})]^T \quad (3)$$

where  $\vec{x} = [x_1, x_2, \dots, x_n]^T$  is the vector of decision variables.  $\square$

**Definition 2 (Pareto Optimality):** A point  $\vec{x}^* \in \Omega$  is Pareto optimal if for every  $\vec{x} \in \Omega$  and  $I = \{1, 2, \dots, k\}$  either,

$$\forall i \in I (f_i(\vec{x}) = f_i(\vec{x}^*)) \quad (4)$$

or, there is at least one  $i \in I$  such that

$$f_i(\vec{x}) > f_i(\vec{x}^*) \quad (5)$$

$\square$

( $\Omega$  is the feasible region). In words, this definition says that  $\vec{x}^*$  is Pareto optimal if there exists no feasible vector  $\vec{x}$  which would decrease some criterion without causing a simultaneous increase in at least one other criterion.

**Definition 3 (Pareto Dominance):** A vector  $\vec{u} = (u_1, \dots, u_k)$  is said to dominate  $\vec{v} = (v_1, \dots, v_k)$  (denoted by  $\vec{u} \preceq \vec{v}$ ) if and only if  $u$  is partially less than  $v$ , i.e.,  $\forall i \in \{1, \dots, k\}, u_i \leq v_i \wedge \exists i \in \{1, \dots, k\} : u_i < v_i$ .  $\square$

**Definition 4 (Pareto Optimal Set):** For a given MOP  $\vec{f}(x)$ , the Pareto optimal set ( $\mathcal{P}^*$ ) is defined as:

$$\mathcal{P}^* := \{x \in \Omega \mid \neg \exists x' \in \Omega \vec{f}(x') \preceq \vec{f}(x)\}. \quad (6)$$

$\square$

**Definition 5 (Pareto Front):** For a given MOP  $\vec{f}(x)$  and Pareto optimal set  $\mathcal{P}^*$ , the Pareto front ( $\mathcal{PF}^*$ ) is defined as:

$$\mathcal{PF}^* := \{\vec{u} = \vec{f} = (f_1(x), \dots, f_k(x)) \mid x \in \mathcal{P}^*\}. \quad (7)$$

$\square$

## III. NOTIONS OF CULTURAL ALGORITHMS

Cultural algorithms were developed by Robert G. Reynolds as a complement to the metaphor adopted by evolutionary algorithms, which was mainly focused on genetic concepts and on the natural selection mechanism [18]. Cultural algorithms are based on some theories proposed in sociology and archaeology to model cultural evolution. Such theories indicate that cultural evolution can be seen as an inheritance process that occurs at two levels: the micro-evolutionary level, and the macro-evolutionary level.

At the micro-evolutionary level, individuals are described in terms of “behavioral traits” (which can be socially acceptable or unacceptable). These behavioral traits are passed from generation to generation using several socially motivated operators. At the macro-evolutionary level, individuals are able to generate

“mappa” [17], or generalized descriptions of their experiences. Individual mappa can be merged and modified to form “group mappa” using a set of generic or problem specific operators. Both levels share a communication link.

The micro-evolutionary level refers to the knowledge acquired by individuals through generations which, once encoded and stored is used to guide the behavior of the individuals that belong to a certain population [17], [7]. Reynolds attempts to capture this double inheritance phenomenon in the cultural algorithm [18]. The goal is to increase the learning or convergence rates of the algorithm as to provide a better response to a large number of problems [10].

Cultural algorithms operate on two spaces. First, they operate on the population space as any other evolutionary computation technique in which a set of individuals (called population) is adopted. Each individual has a set of features independent from each other which allows us to determine its fitness. Through time, such individuals can be replaced by some of its descendants, obtained after applying a set of operators to the population.

The second space is the belief space, where the knowledge acquired by the individuals along the evolutionary process is stored. The information contained in this space must be accessible to any individual, so that it can use it to modify its behavior.

To unify both spaces, a communication protocol is established such that it dictates rules regarding the type of information to be exchanged between these two spaces. For example, to update the belief space, the individual experiences of a select set of individuals are incorporated. This select group of individuals is obtained with the function *acceptance* which is applied to the entire population. On the other hand, the operators that modify the population (i.e., recombination and mutation) and the selection operator are modified by the function *influence*. This function acts in such a way that the individuals resulting from the application of the operators tend to approach the desirable behavior while staying away from undesirable behaviors. Such desirable and undesirable behaviors are defined in terms of the information stored in the belief space. These two functions are used to establish the communication between the two spaces (i.e., population and belief). The interactions between these two spaces can be appreciated in Figure 1 [19].

Following this model, we have designed a cultural algorithm for multiobjective optimization, trying to take advantage of its main features.

#### IV. DESCRIPTION OF OUR APPROACH

We propose the use of a cultural algorithm combined with evolutionary programming (*Cultural Algorithm with Evolutionary Programming* [1], or CAEP). The pseudocode of our approach is shown in Algorithm 1. Here, we can clearly see the similarities of our approach with traditional evolutionary programming [8], and we also include the steps where the belief space is incorporated.

The problems that we are interested in solving have  $n$  decision variables and  $k$  objective functions. The population consists of a set of individuals, each of which repre-

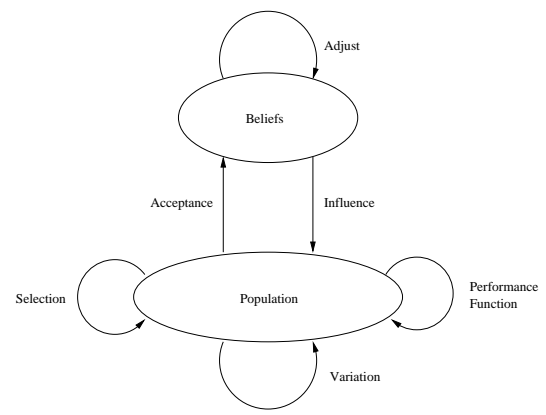


Fig. 1. Spaces of a cultural algorithm.

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#### Algorithm 1 Basic structure of the cultural algorithm for multiobjective optimization.

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Generate the initial population of size  $p$ 
Evaluate the initial population
Initialize the belief space
Repeat
    Apply mutation
        to generate  $p$  offspring (we now
        have  $2p$  individuals in the population)
    Evaluate each offspring
    Perform binary tournaments, randomly
        choosing  $c$  contenders for each
        individual. The decisions taken
        in the tournaments will be
        influenced by the information
        stored in the belief space.
    Select the  $p$  individuals with
        the largest number of victories
        in the tournaments, to produce the
        population of the following generation
    Add the new nondominated individuals
        to the external memory (file)
    Update belief space with the individuals
        added to the external memory
While stopping condition is not met

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sents a possible solution to the problem. Each individual contains the  $n$  decision variables of the problem to be solved. The population is initialized with  $p$  individuals randomly generated using a uniform distribution within the allowable range for each decision variable. The external memory indicated in the Algorithm 1 is an external file (or secondary population) where the nondominated individuals found along the evolutionary are stored. The final contents of this file is the set of solutions presented by the algorithm to the user. This external memory has a maximum size  $q$  that corresponds to the number of nondominated solutions that we aim to obtain. Next, we detail the structure of the belief space adopted, together with the remaining steps of the algorithm.

##### A. Structure of the Belief Space

The belief space consists of two parts: the phenotypic normative part and a grid which is used to emphasize the

$l_{f1}$	$u_{f1}$	$l_{f2}$	$u_{f2}$	$\dots$	$l_{fk}$	$u_{fk}$
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Fig. 2. Phenotypic normative part

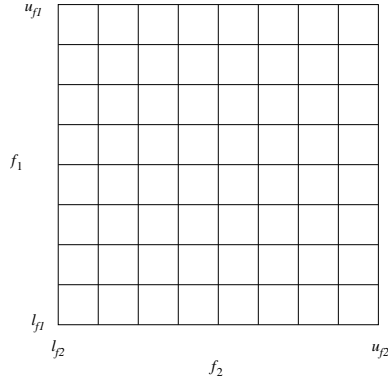


Fig. 3. Grid in the belief space for a problem with two objective functions. In this case,  $s_1 = s_2 = 8$  (eight sub-intervals in each dimension).

generation of nondominated solutions that are uniformly distributed along the Pareto front. This grid is a variation of the adaptive grid proposed by Knowles and Corne [14].

The phenotypic normative part contains only the lower and upper bounds,  $l_{fi}$  and  $u_{fi}$ , of the intervals for each objective function ( $i = 1, \dots, k$ ) within which the grid will be built. This grid will be used to place each nondominated solution in some sort of coordinate system where the values of the objective functions are used to place each solution (see Figure 2).

Once we have these intervals, we only need to know the number of identical sub-divisions to apply to each of them ( $s_i$ , with  $i = 1, \dots, k$ ), so that we can build the grid in phenotypic space. An example of the grid adopted is shown in Figure 3.

As a result, we will have  $s_1 s_2 \dots s_k$  cells, which will all have the same dimensions. The values  $s_i$  are input parameters required by the algorithm. For each cell, we store the count of the number of nondominated individuals stored in the secondary memory that are contained within. This is useful to obtain an appropriate distribution of the nondominated solutions, avoiding that they all cluster together around a certain portion of the Pareto front.

## B. Initialization of the Belief Space

In order to initialize the belief space is necessary to have an initial population, because we will use the nondominated individuals from that population (it can be proved that in any population of size greater than zero there is at least one nondominated individual [23]).

### B.1 Initialization of the Phenotypic Normative Part

The initialization of the phenotypic normative part of the belief space consists of finding the extrema of each objective function for the nondominated individuals of the initial population. These extrema are stored in  $l_{fi}$  and  $u_{fi}$ , so that we can place the grid in the region where the nondominated individuals known so far are located.

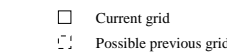


Fig. 4. The update of the phenotypic normative part of the belief space aims to contain the entire Pareto front known so far within the grid.

## B.2 Initialization of the Grid

The grid is created taking as intervals the values stored in the phenotypic normative part, and we use the input parameters  $s_i$  to divide such grid. The counters of the nondominated individuals contained within each cell are initialized to zero.

### C. Updating Belief Space

The belief space grid is updated at each generation, whereas the phenotypic normative part is updated at each  $g_{normative}$  generations, where  $g_{normative}$  is a parameter defined by the user.

#### C.1 Updating the Grid

In order to update the grid, we simply increment the counters of the nondominated individuals by the number of individuals added to the external memory during the current generation. The update of the grid is very simple, and that is the reason why we do it at every generation. For the update of this part of the belief space, the acceptance function uses the population of the external memory and only chooses the new individuals within that population.

#### C.2 Updating the Phenotypic Normative Part

The update of the phenotypic normative part is not done at each generation, because it involves the rebuilding of the grid and, therefore, it is a process that could seriously affect the computational efficiency of the algorithm. For this update, we also use the population contained in the secondary memory.

Again, as in the initialization of this phenotypic normative part, we need to identify the extrema for each objective function contained in the external memory. These values are stored in  $l_{fi}$  and  $u_{fi}$  for  $i = 1, \dots, k$ . With the new values, we build the grid, which covers the entire Pareto front known so far (see Figure 4) and that probably has limits beyond the boundaries of the previous grid.

After initializing the grid, all the counters are set to zero and it is then necessary to add all the individuals from the external memory to the counter of their corresponding cell, so that the belief space is ready again to be used.

#### D. Mutation

The information stored in the belief space belongs to the phenotypic space of our problem. Therefore, it is difficult to use it to self-adapt the mutation operator (since mutation operates on genotypic space). Given this inconvenience, we leave the mutation parameters as inputs for the algorithm, which must be provided by the user (i.e., no self-adaptation mechanism is adopted).

The Gaussian mutation operator adopted in our algorithm is based on the following expression:

$$x'_i = x_i + N(0, \sigma)$$

where:  $x_i$  is the  $i$ -th variable of individual  $x$ ,  $x'_i$  is the  $i$ -th variable of the new individual  $x'$  obtained after applying mutation, and  $N(\mu, \sigma)$  is a random variable with a normal distribution that has a mean  $\mu$  and a standard deviation  $\sigma$ . In our case,  $\mu$  will always be zero, and  $\sigma$  is a parameter provided by the user. Mutation is then applied for  $i = 1, \dots, n$ , and it operates on the main population. Therefore, at the end of the process, we will have a population of size  $2p$ .

#### E. Tournament Selection

Tournament selection is performed considering a main population of size  $2p$ . Each individual is confronted against other  $c$  individuals which are randomly chosen from the main population. Tournament rules are the following:

1. If an individual dominates his contender, then the dominating individual wins.
2. If the individuals confronted are non-comparable, or if the values of their objective functions are the same, then:
  - (a) If both lie within the grid of the belief space, then the individual located in the less populated cell (according to the cell counters) wins.
  - (b) If one of the individuals lies outside the grid, then it automatically wins.

The first rule should be straightforward, since we are simply giving preference to nondominated individuals so that we can approach the Pareto front. In the second point, we appreciate the influence of the belief space in the decisions made during the tournament. The first subpart of the second point aims to distribute solutions in a uniform way among the cells, so that we can produce a better distributed Pareto front. Finally, if we have the case indicated in the second subpart of the second point, it means that we have found a solution that is beyond the Pareto front known so far. Since this solution will generate a new segment of the Pareto front, it is important to keep it. Figure 5 shows the case in which an individual lies outside the grid.

Once the tournaments finish, we select the individuals with the largest number of victories to be part of the following generation. We can see that these tournaments are similar to those of the NPGA [11], where each tournament competitor is compared with respect to random sample of the population. However, in the NPGA we apply Pareto ranking with respect to the sample randomly

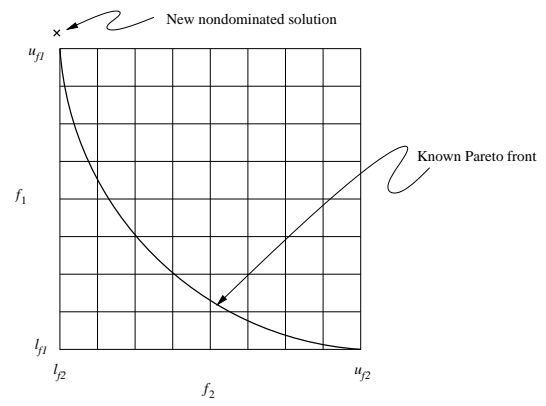


Fig. 5. Tournament in the case in which an individual lies outside the grid. The point found is a new extrema of the Pareto front known so far and it is therefore important to keep it.

selected, whereas in our algorithm we compare individually, counting the number of victories (as it is normally done in evolutionary programming).

#### F. Adding Individuals to the External Memory

The external memory must contain only nondominated individuals, without repetition. To add such individuals to the external memory, we use the following rules:

1. If the individual that we want to add is dominated by another individual contained in the external memory, then the new individual is not added.
2. If the individual that we want to add dominates some individual contained in the external memory, then it is introduced in its place, but it is still compared with respect to everybody else. If the same individual recently added dominates another one, then the dominated individual is removed from the external memory.
3. If the individual that we want to add is not dominated and it does not dominate any other individual in the external memory, and if the number of individuals in the external memory is less than the maximum allowable value  $q$ , then we add the new individual at the end.
4. If the individual that we want to add is not dominated and it does not dominate any other individual in the external memory, but the external memory is already full, then we locate an individual whose cell contains more individuals than the cell to which the new individual would belong (in case it is added). Then, we replace the old individual by the new one. This will motivate a better distribution of nondominated solutions along the Pareto front.

### V. COMPARISON OF RESULTS

In order to validate our proposed approach, we used a set of test functions which have commonly been adopted as a benchmark in the specialized literature [3]. The first test function, **MOP1**, was proposed by Schaffer [21]; our second test function, **MOP2**, was proposed by Fonseca [9]; our third test function **MOP3** was proposed by Poloni [16]. **MOP4** was proposed by Kursawe [15]; Finally, **MOP6** was designed following Deb's methodology [5]. All these functions have variable

degrees of difficulty. There are problems with convex and concave Pareto fronts, continuous and disconnected, and problems with two and with three objective functions. The mathematical description of these problems is provided next:

**MOP1.** Minimize:

$$\vec{f}(x) = (f_1(x), f_2(x))$$

where:

$$\begin{aligned} f_1(x) &= x^2 \\ f_2(x) &= (x-2)^2 \end{aligned}$$

and  $-10^5 \leq x \leq 10^5$ .

**MOP2.** Minimize:

$$\vec{f}(\vec{x}) = (f_1(\vec{x}), f_2(\vec{x}))$$

where:

$$\begin{aligned} f_1(\vec{x}) &= 1 - \exp\left(-\sum_{i=1}^n \left(x_i - \frac{1}{\sqrt{n}}\right)^2\right) \\ f_2(\vec{x}) &= 1 - \exp\left(-\sum_{i=1}^n \left(x_i + \frac{1}{\sqrt{n}}\right)^2\right) \end{aligned}$$

and  $-4 \leq x_i \leq 4$  ( $i = 1, 2, 3$ ).

**MOP3.** Maximize:

$$\vec{f}(x, y) = (f_1(x, y), f_2(x, y))$$

where:

$$\begin{aligned} f_1(x, y) &= -[1 + (A_1 - B_1)^2 + (A_2 - B_2)^2] \\ f_2(x, y) &= -[(x+3)^2 + (y+1)^2] \end{aligned}$$

with:

$$\begin{aligned} A_1 &= 0.5 \sin 1 - 2 \cos 1 + \sin 2 - 1.5 \cos 2 \\ A_2 &= 1.5 \sin 1 - \cos 1 + 2 \sin 2 - 0.5 \cos 2 \\ B_1 &= 0.5 \sin x - 2 \cos x + \sin y - 1.5 \cos y \\ B_2 &= 1.5 \sin x - \cos x + 2 \sin y - 0.5 \cos y \end{aligned}$$

and  $-\pi \leq x, y \leq \pi$ .

**MOP4.** Minimize:

$$\vec{f}(\vec{x}) = (f_1(\vec{x}), f_2(\vec{x}))$$

where:

$$\begin{aligned} f_1(\vec{x}) &= \sum_{i=1}^{n-1} \left(-10e^{(-0.2)\sqrt{x_i^2 + x_{i+1}^2}}\right) \\ f_2(\vec{x}) &= \sum_{i=1}^n (|x_i|^a + 5 \sin(x_i^b)) \end{aligned}$$

with:

$$\begin{aligned} a &= 0.8 \\ b &= 3 \end{aligned}$$

and  $-5 \leq x_i \leq 5$  ( $i = 1, 2, 3$ ).

**MOP6.** Minimize:

$$\vec{f}(x, y) = (f_1(x, y), f_2(x, y))$$

where:

$$\begin{aligned} f_1(x, y) &= x \\ f_2(x, y) &= (1 + 10y) \left[ 1 - \left( \frac{x}{1 + 10y} \right)^\alpha - \frac{x}{1 + 10y} \sin(2\pi qx) \right] \end{aligned}$$

with:  $q = 4, \alpha = 2, y \ 0 \leq x, y \leq 1$ .

The parameters adopted by our CAEP are the following:  $p = 6, q = 100, G_{max} = 35,000, g_{normative} = 20, s_i = 10$  ( $i = 1, \dots, k$ ),  $c = \frac{p}{2} = 5, \sigma = 1$ . The total number of objective function evaluations during one execution of our algorithm is computed using  $p(G_{max} + 1)$ . For the examples used in this paper, we performed 210,006 evaluations per run.

We compared CAEP with respect to the NSGA-II [6], which used the following parameters: population size = 100, maximum number of generations = 2100, crossover rate = 0.9, mutation rate =  $\frac{1}{n}$ , SBX parameter = 10 y mutation parameter = 100. These values have been suggested by the author of the algorithm, except for the maximum number of generations, which was chosen in such a way that the total number of evaluations performed by the NSGA-II was approximately the same as CAEP. The NSGA-II performed 210,100 objective function evaluations per run.

Table I shows the statistics of each of the three metrics adopted to allow a quantitative comparison: error ratio (ER) [23], generational distance (GD) [23] and spacing (SP) [22] for ten independent runs of each algorithm.

Figure 6 shows the average behavior of each of the two algorithms compared (CAEP y NSGA-II) for the first test function. Based on the results shown in Table I we can see that, in this case, CAEP obtained better results for the three metrics adopted.

Figure 7 shows the average behavior of each of the two algorithms compared in **MOP2**. In this case, our approach had a high error ratio (approximately 50%). With respect to generational distance, both algorithms had low values. With respect to spacing, the NSGA-II had values slightly higher than those of CAEP.

Figure 8 shows the average behavior of each of the two algorithms compared in **MOP3**. In this case, the NSGA-II had a lower generational distance than our approach, but a higher error ratio. This problem has a disconnected Pareto front and, therefore, spacing does not correctly reflect the performance of an algorithm. With respect to spacing, the NSGA-II had better values than CAEP.

Figure 9 shows the average behavior of the two algorithms in **MOP4**. In this case, all the metrics indicate better values for the NSGA-II, but our algorithm covers a region of the Pareto front that the NSGA-II can never cover, as Figure 9 indicates.

Figure 10 shows the average behavior of the two algorithms in **MOP6**. In this case, both algorithms have

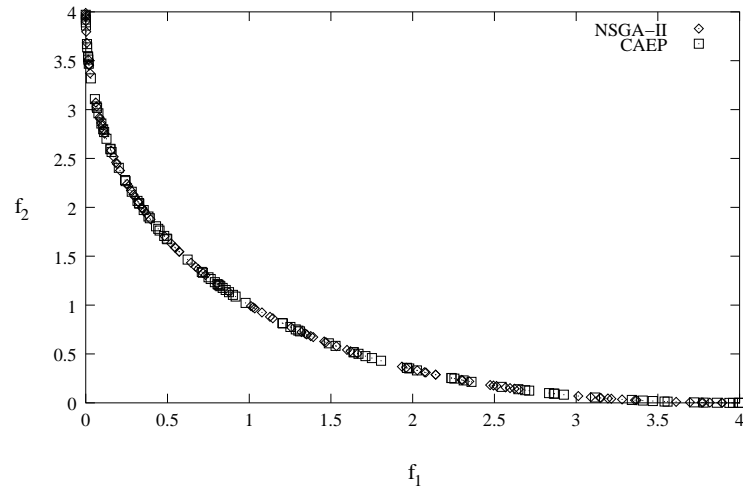


Fig. 6. Comparison of results between our algorithm (CAEP) and the NSGA-II for **MOP1**.

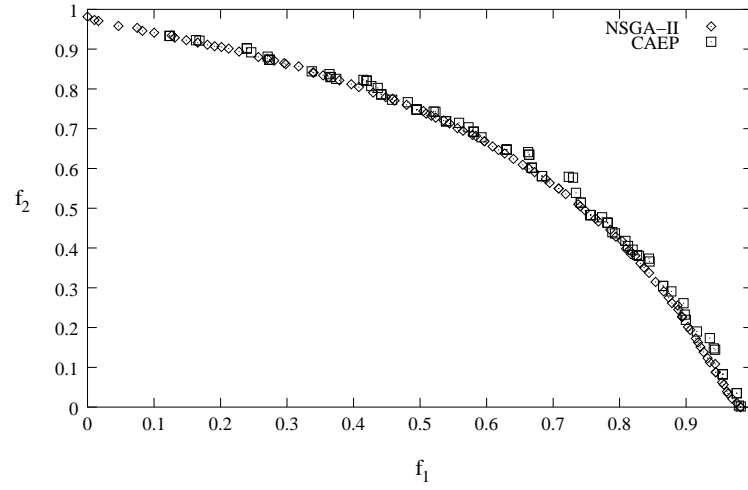


Fig. 7. Comparison of results between our algorithm (CAEP) and the NSGA-II for **MOP2**.

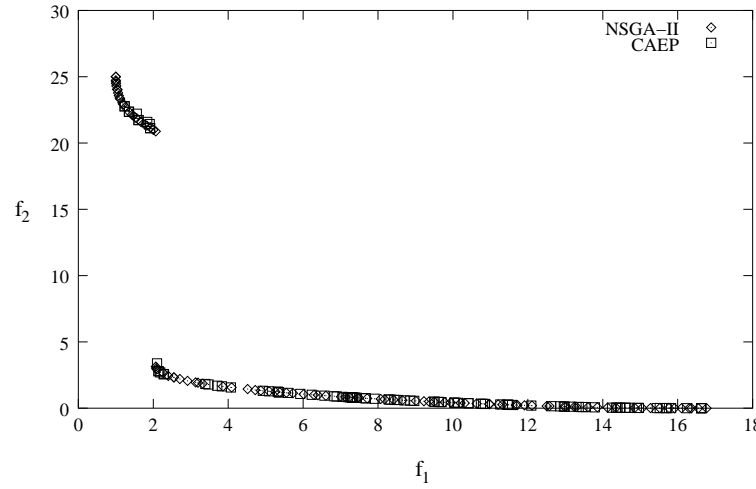


Fig. 8. Comparison of results between our algorithm (CAEP) and the NSGA-II for **MOP3**.

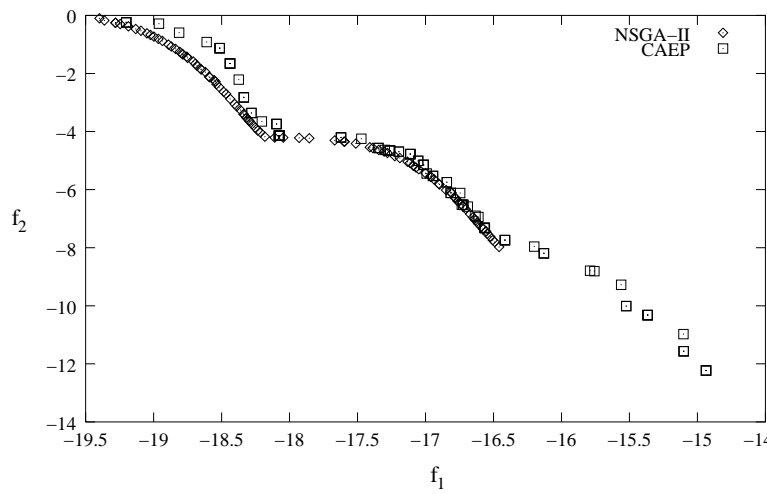


Fig. 9. Comparison of results between our algorithm (CAEP) and the NSGA-II for **MOP4**.

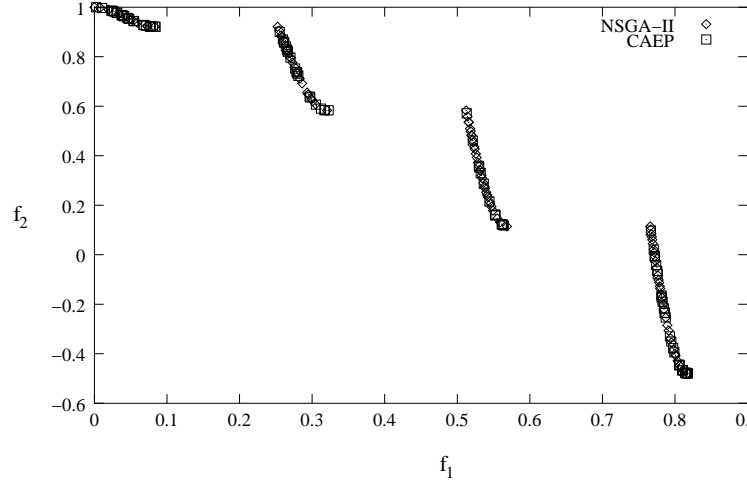


Fig. 10. Comparison of results between our algorithm (CAEP) and the NSGA-II for **MOP6**.

very low error ratio values (the NSGA-II almost always produces 0%), but our approach has a lower generational distance than the NSGA-II. Regarding spacing, we again have values slightly better for the NSGA-II, but this is caused by the disconnected Pareto front of this problem.

In summary, our results indicate that CAEP has a competitive behavior with respect to the NSGA-II, which is representative of the state-of-the-art in evolutionary multiobjective optimization.

## VI. CONCLUSIONS AND FUTURE WORK

In this paper we have introduced the first proposal to use cultural algorithms for multiobjective optimization incorporating Pareto-based selection. We have shown how adding a belief space to an evolutionary programming algorithm can be beneficial and represents a viable alternative to solve multiobjective problems.

The small comparative study presented in this paper validates the viability of our proposal. However, we obviously still need to do a more in-depth study in which other evolutionary multiobjective optimization techniques and more test functions are involved. Additionally, there are still a few limitations of CAEP that require a more careful analysis. For example, CAEP tends to lose diversity very quickly in some cases. In order to deal with this prob-

lem, we are currently designing an alternative scheme to maintain diversity which emphasizes efficiency (computationally speaking). We are also interested in improving the mechanism adopted to generate a uniform distribution of nondominated solutions along the Pareto front. This is motivated by the fact that our current mechanism still has some flaws and sometimes the distribution produced by our algorithm is not as good as expected.

## ACKNOWLEDGMENTS

The first author acknowledges support from the Mexican Consejo Nacional de Ciencia y Tecnología (CONACyT) through project number 34201-A. The second author acknowledges support from CONACyT through a scholarship to pursue graduate studies at CINVESTAV-IPN's Electrical Engineering Department (Computer Science Section).

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TABLE I  
COMPARISON OF RESULTS.

MOP		ER		GD		SP	
		CAEP	NSGA-II	CAEP	NSGA-II	CAEP	NSGA-II
1	Average	0.003	0.008059	0.000946	0.061548	0.037342	0.058298
	Std. Dev.	0.006749	0.005818	0.000046	0.002711	0.002674	0.009385
	Min.	0	0	0.000899	0.058940	0.034057	0.043401
	Max.	0.02	0.017857	0.001045	0.067989	0.042604	0.075381
2	Average	0.911	0.574	0.001103	0.000280	0.010931	0.007241
	Std. Dev.	0.046774	0.034059	0.000171	0.000034	0.001040	0.000741
	Min.	0.84	0.53	0.000934	0.000230	0.009493	0.006015
	Max.	0.98	0.63	0.001514	0.000349	0.012157	0.008287
3	Average	0.149	0.209	0.015321	0.001941	0.607229	0.092216
	Std. Dev.	0.069194	0.041218	0.026480	0.000078	1.025953	0.008415
	Min.	0.06	0.15	0.001942	0.001822	0.065285	0.081569
	Max.	0.28	0.28	0.076256	0.002107	2.80319	0.106568
4	Average	0.876	0.144	0.022522	0.002892	0.139053	0.038378
	Std. Dev.	0.101017	0.049035	0.008002	0.000203	0.049507	0.003837
	Min.	0.64	0.05	0.013604	0.002525	0.094707	0.031393
	Max.	0.97	0.21	0.037278	0.003199	0.242052	0.044254
6	Average	0.028	0	0.000259	0.000337	0.014476	0.008266
	Std. Dev.	0.018135	0	0.000028	0.000013	0.005168	0.000918
	Min.	0	0	0.000220	0.000318	0.009296	0.006851
	Max.	0.06	0	0.000302	0.000355	0.022920	0.010127

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