

# Multiobjective Optimization using Genetic Programming: Reducing Selection Pressure by Approximate Dominance

Ayman Elkasaby, Akram Salah and Ehab Elfeky  
*Faculty of Computers and Information, Cairo University, Giza, Cairo, Egypt*

**Keywords:** Genetic Programming, Multiobjective Optimization, Epsilon Dominance, Evolutionary Algorithms.

**Abstract:** Multi-objective optimization is currently an active area of research, due to the difficulty of obtaining diverse and high-quality solutions quickly. Focusing on the diversity or quality aspect means deterioration of the other, while optimizing both results in impractically long computational times. This gives rise to approximate measures, which relax the constraints and manage to obtain good-enough results in suitable running times. One such measure, epsilon-dominance, relaxes the criteria by which a solution dominates another. Combining this measure with genetic programming, an evolutionary algorithm that is flexible and can solve sophisticated problems, makes it potentially useful in solving difficult optimization problems. Preliminary results on small problems prove the efficacy of the method and suggest its potential on problems with more objectives.

## 1 INTRODUCTION

Historically, in order to solve optimization problems, classical search methods were traditionally used. In every iteration, a single solution was modified in order to produce better solutions. However, this point-by-point approach was overshadowed by the introduction of evolutionary algorithms (EAs). These algorithms use the concepts of evolution and natural selection in optimization. Using populations of individual solutions, EAs try to capture multiple optimal solutions for problems lacking one global optimal solution.

Some optimization, for example industrial, problems have multiple objectives that need to be optimized in the same time, which poses extra difficulties for algorithms that try to solve these problems. Two main solutions have usually been followed to reduce the complexities:

- Reducing the number of objectives during the search process or a posteriori during the decision making process. This approach tries to identify non-conflicting objectives and discards them.

- Propose a preference relation that induces a finer order on the objective space.

If the aforementioned solutions fail to reduce multi-objective optimization problems' complexity, then the main difficulty facing EAs is incomparable

solutions. Incomparable solutions happen in the following case. When one solution optimizes one (or more) objective better than a second solution, but the second solution optimizes another (or more) different objective better than the first one.

If we divide the search space into regions based on how well each solution optimizes each objective, and assuming no bias towards any region, the probability of a solution falling into any of these regions is proportional to the volume of this region divided by the volume of the entire solution set. As the number of objectives increases, the number of regions increases, and the probability that a solution will fall into a region where one solution optimizes all objectives efficiently is reduced significantly.

Problems with a large number of objectives, although apparently similar to problems with less number of objectives, can't be solved efficiently using the same methods used for fewer objectives. They are computationally more intensive, and visualizing their solutions becomes harder as more objectives are added. To avoid these complexities, some approximate measures are used to obtain good-enough results of the problem. Epsilon dominance, notated as  $\epsilon$ -dominance from now on, is one of these approximate measures (Laumanns, et al., 2002).

In this paper, genetic programming, a flexible and powerful type of evolutionary algorithms (EAs), is used in order to solve optimization problems

approximately using  $\epsilon$ -dominance. We call this method  $\epsilon$ -GP. Genetic programming, up to our knowledge, has not been used before to solve any optimization problem with an approximate measure.  $\epsilon$ -GP is compared to regular genetic programming (Koza, 1992) in regards to speed, efficiency, and diversity, and it gives promising results.

The paper is structured as follows. **Section 2** explains related work in the field of evolutionary algorithms. Afterwards, in **Section 3 and 4**, some background information is given about optimization and genetic programming, respectively. An outline and pseudocode of  $\epsilon$ -GP are given in **Section 5**, while **Section 6** deals with the experimentation and results. Finally, **Section 7** contains a conclusion of the paper and explains future work.

## 2 RELATED WORK

Evolutionary algorithms have long been successful in solving MOPs. Schaffer (Schaffer, 1985) started the movement of EAs solving MOPs by introducing a vector-evaluated genetic algorithm (VEGA) that finds a set of nondominated solutions.

Afterwards, the first generation of Multi-Objective Optimization Evolutionary Algorithms (MOEAs) started in the early 1990s by using Pareto ranking and fitness sharing. This generation consisted of the multi-objective genetic algorithm (MOGA) (Fonseca & Fleming, 1993), the niched Pareto genetic algorithm (NPGA) (Abido, 2003) which is the first algorithm to use tournament selection, and the nondominated sorting algorithm (NSGA) (Srinivas & Deb, 1994).

The second generation of MOEAs, which emerged in the late 1990s and early 2000s, introduced the concept of elitism (keeping a record of the best-so-far solutions). It includes the strength Pareto evolutionary algorithm (SPEA) (Zitzler & Thiele, 1999) and its improved version (SPEA-2) which adds a fitness assignment technique, a nearest neighbor density estimation, and a preservation truncation method (Zitzler, et al., 2001); the Pareto archived evolution strategy (PAES) (Knowles & Corne, 2000); the Pareto envelope based algorithm (PESA) (Corne, et al., 2000) and its improved version PESA-II (Corne, et al., 2001); and an improved version of NSGA (NSGA-II) which splits the pool of individuals into different fronts according to their dominance and adds a crowding measure to preserve diversity (Deb, et al., 2002).

NSGA-II is one of the most popular algorithms in the literature and is usually considered a benchmark for many new algorithms. This is

because it is very quick in obtaining solutions. It also yields very efficient results. Although originally made for problems with smaller number of objectives, NSGA-II has shown to be somewhat successful over the years in solving some problems with more objectives as well.

## 3 OPTIMIZATION

An optimization problem is a problem where the goal is to find the best solution from all feasible solutions for a specific objective function. However, many of these problems (those that have more than one objective) exist in a setting that cannot be expressed using a single function, as different objectives are usually not measured using the same metrics.

Furthermore, a multi-objective optimization problem is defined as simultaneously optimizing

$$F(x) = \min(f_1(x), \dots, f_k(x)), \quad (1)$$

*subject to*  $x \in \bar{X}$ ,

by changing  $n$  decision variables, subject to some constraints that define the universe  $\bar{X}$ .

In other words, a multi-objective optimization solution optimizes the components of  $F(x)$  where  $x$  is an  $n$ -dimensional decision variable vector  $x = (x_1, \dots, x_n)$  from some universe  $\bar{X}$ . Thus, the problem consists of  $k$  objectives reflected in the  $k$  objective functions, a number of constraints on the objective functions reflected on the feasible set of decision vectors  $\bar{X}$ , and  $n$  decision variables.

In the case of optimizing multiple objectives, it is usually impossible to find a single solution that optimizes all of the objectives at the same time. This gives rise to the definition of nondominated solutions (also called Pareto-optimal solutions), which are solutions that optimize some objectives but are not worse than other solutions in the rest of the objectives. The Pareto front is the visualization of all these solutions on the search space. Since these solutions are nondominated, no one solution exists that can be said to be better than the other; all of them are presented to the decision maker as a set of solutions called the Pareto optimal set.

Multi-objective optimizers usually have to conform to a few properties; namely, they should present solutions that are close to the Pareto front as possible. They should also present different, diverse solutions to the decision maker that show the

different tradeoffs with respect to each objective. Optimizers also need to present the best few, which means that overwhelming the decision maker by presenting too many solutions is not preferred.

### 3.1 More Objectives

Optimization problems that have more than 3 objectives are named *many*-objective optimization problems, and problems with 2 or 3 objectives are named *multi*-objective optimization problems. In (Khare, et al., 2003), it was found after testing 3 MOEAs from the 2<sup>nd</sup> generation of MOEAs (NSGA-II, SPEA2, PESA) that these algorithms showed vulnerability on problems with a larger number of objectives.

The main difficulties with many-objective optimization problems are visualization, how to handle high dimensionality and the exponential number of points needed to represent the Pareto front, the greater proportion of nondominated solutions, and stagnation of search due to larger number of incomparable solutions. Our work tackles the latter two difficulties by changing the definition of dominance to an approximate one, easing the criteria of acceptance of nondominated solutions.

### 3.2 Dominance

Multi-objective optimization algorithms insisting on both diversity and convergence to the Pareto front face Pareto sets of substantial sizes, need huge computation time, and are forced to present very large solutions to the decision maker. These issues effectively make them useless until further analysis, because speed and presenting few solutions are very important to decision makers.

$\epsilon$ -dominance (Laumanns, et al., 2002) tries to fix these problems by quickly searching for solutions that are good enough, diverse, and few in number. It approximates domination in the Pareto set by relaxing the strict definitions of dominance and considering individuals to  $\epsilon$ -dominate other individuals, whereas previously they would have been nondominated to each other.

In Figure 1, a visual comparison between  $\epsilon$ -dominance and regular dominance is shown (Laumanns, et al., 2002).

## 4 GENETIC PROGRAMMING

Genetic programming (GP) is one type of evolutionary algorithms. Its main characteristic is

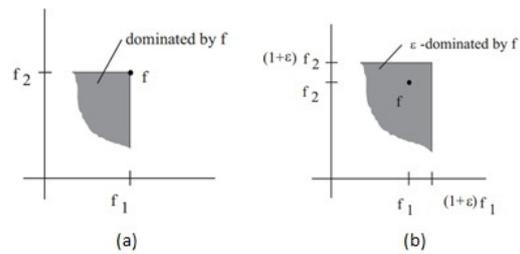


Figure 1: Differences between (a) regular and (b)  $\epsilon$  - dominance.

that it represents solutions as programs (Koza, 1992). This representation scheme is the main difference between genetic algorithms and genetic programming. Each solution (program) is judged based on its ability to solve the problem, using a mathematical function, the fitness function. Each program, or solution, is represented using a decision tree. GP evolves a population of programs by selecting some candidates that score high on the fitness function and using regular evolutionary variation operators on them (mutation, crossover, and reproduction). New populations are created from these outputs until any specific termination criterion is met.

We use strongly-typed genetic programming (STGP) in this paper, which is one of many enhanced versions of GP. STGP makes GP more flexible, explicitly defining allowed data types beforehand instead of limiting it to only one data type. Genetic programming, and STGP specifically, consists of the following.

1) Representation: individuals are represented as decision trees, but unlike usual GP (Koza, 1992), STGP doesn't limit variables, constants, arguments for functions, and values returned from functions to be of the same data type. We only need to specify the data types beforehand. Additionally, to ensure consistency, the root node of the tree must return a value of the type specified by the problem definition and each nonroot node has to return a value of the type required by its parent node as an argument.

2) Fitness function: scores how well a specific execution matches expected results.

3) Initialization: there are two main methods to initialize a population: *full* and *grow*. Koza (Koza, 1992) recommended using a ramped half-and-half approach, combining the two methods equally.

4) Genetic operators: crossover and mutation.

5) Parameters: maximum tree depth, maximum initial tree depth, max mutation tree depth, population size, and termination criteria.

## 5 OUR PROPOSED METHOD

Our algorithm,  $\epsilon$ -GP, has three main characteristics. First, the performance of our algorithm, and of any general  $\epsilon$ -dominance-based MOEA, depends on the value of  $\epsilon$ , which is either user defined or computed from the number of solutions required. Bigger  $\epsilon$  values mean quicker computation of solutions, while smaller values mean solutions that have more quality. Although the value of  $\epsilon$  doesn't have to be constant for each objective, we make it constant across all objectives in our method for ease of use and for quicker computations.

Second,  $\epsilon$ -GP comprises two storage locations for solutions:

- An archive that ensures elitism by keeping the best solutions so far and removing solutions iff other better solutions are found. We choose to give this archive a fixed size for several reasons. One is to limit computation time and to protect the decision maker from receiving a big number of nondominated solutions that are incomparable. Finally, due to  $\epsilon$ -MOEA performing well on all test instances in (Li, et al., 2013), while suffering from archive size instability, we will stabilize and fix the size.
- A population that stores the current generation; this current generation can have worse solutions than a previous generation. This ensures diversity and keeps us from falling into local minima.

Third, crossover is always between a solution from the current generation population and a solution from the archive. This guarantees both elitism and diversity. Offspring from crossover are embedded into the archive if the criteria of acceptance (to dominate another solution) are met. They are automatically inserted into the next generation population as well.

To our knowledge,  $\epsilon$ -GP is the first algorithm to combine genetic programming with an approximate measure, e.g.,  $\epsilon$ -dominance.  $\epsilon$ -GP uses its approximation capability to make selection easier between points by reducing competition and tolerating a certain additive factor ( $\epsilon$ ) when calculating dominance. Selection is the most computation-intensive regular task in Many-O algorithms, and this is why  $\epsilon$ -GP is considered useful.

The initial random generation of the population and archive in our algorithm is done using the ramped half-and-half method discussed earlier. The user inputs in  $\epsilon$ -GP are the number of runs, the population size (`pop_size`), the probabilities  $G_0$ ,  $Pr$ ,

$P_c$  (respectively, probability of a binary or unary genetic operator, probability of reproduction or mutation, and probability of crossover).

The pseudocode of  $\epsilon$ -GP is as follows:

```

for (i = 0; i < number_of_runs; i++) {
    set gen, score to 0; //generation
    number

    generate pop[gen], archive randomly;

    while (score <= minimum_threshold &&
    generation < max_generations) {

        evaluate fitness of pop[gen];

        sort individuals in archive and
        pop; // this is where  $\epsilon$ -dominance is
        used

        for (j = 0; j < pop_size; j++) {
            if (random(0,1) >=  $G_0$ ) {
                if (random(0,1) >=  $Pr$ ) {
                    reproduce (copy) individual;
                }
            }
            else {
                mutate individual;
            }
            put individual into pop[gen+1];
        }
        else {
            select two individuals from
            archive and pop;
            if(random(0,1) >=  $P_c$ ) {
                crossover the individuals;
            }
            else {
                reproduce both individuals;
            }
            j++; //because we insert 2, not
            1, individuals
            put individuals to pop[gen+1];
        }
        if (individuals(s) >
        archive.worse_result) {
            put individual (s) in archive;
            score = fitness of archive;
        }
    }
}

```

```

    }
    generation ++;
}
set result[i] to score;
}

```

## 6 EXPERIMENTATION

To measure the performance of our algorithm, it was tested on a basic genetic programming problem: the ant trail problem (Koza, 1992). Two variants of this problem are tested; namely, the Santa Fe Trail problem and the Los Altos Trail problem. The study used the MOEA Framework, version 2.8, available from <http://www.moeaframework.org/>. Koza’s GP (Koza, 1992) was used as a reference for comparison in terms of speed and efficiency. We tested the reference against our algorithm with values of  $\epsilon$  of 0.1 and 0.01.

We solved each test problem 30 times with different random seeds. In all runs, no more than 500,000 evaluations were allowed to be made. We used a crossover probability rate of 0.9, with a point mutation rate at 0.01. Population size was set to 500.

Since GP is a stochastic algorithm that is affected by the chosen random seed, it was more suitable to make a stochastic comparison instead of a static comparison with the best absolute values. For this purpose, we analyzed the mean, median, and standard deviation of the 30 independent runs.

The Santa Fe problem results are shown in Table 1, with better results highlighted in bold when applicable. The goal is to capture as much pieces of food as possible, with as little moves as possible. We also take into consideration how quickly a run reaches a suitable result.

Table 1: Santa Fe Trail Results.

Variables	Santa Fe Trail					
	Mean	Median	St. Dev.	Worst	Best	
Moves	Koza	400.533	451	88.1807	494	234
	$\epsilon = 0.1$	<b>368.067</b>	<b>366</b>	89.7241	<b>492</b>	<b>226</b>
	$\epsilon = 0.01$	380.733	390	<b>85.9386</b>	496	230
Food	Koza	80.3	88	10.3629	55	89
	$\epsilon = 0.1$	<b>81.1</b>	<b>89</b>	11.2322	52	89
	$\epsilon = 0.01$	80.1	86.5	<b>10.145</b>	<b>57</b>	89
Time	Koza	64.5333	64	8.91621	<b>79</b>	<b>45</b>
	$\epsilon = 0.1$	<b>58.5</b>	<b>56.5</b>	8.94331	85	46
	$\epsilon = 0.01$	61.7	61	<b>8.78145</b>	81	46

The results show that our algorithm,  $\epsilon$ -GP, has very good performance with regards to all objectives, and runs quickly as well. At both  $\epsilon$

values of 0.1 and 0.01,  $\epsilon$ -GP has a better average runtime compared to Koza’s GP, and at  $\epsilon$  of 0.1, the food gathered by  $\epsilon$ -GP is, on average, more than the food gathered by Koza’s GP.

Next, we test the Los Altos problem, a similar but harder problem with a more complex trail to follow to gather the food.

Table 2: Los Altos Trail.

Variables	Los Altos Trail					
	Mean	Median	St. Dev.	Worst	Best	
Moves	Koza	393.7667	414	80.32635	<b>497</b>	221
	$\epsilon = 0.1$	402.1	412	81.66346	499	<b>132</b>
	$\epsilon = 0.01$	<b>389.667</b>	<b>375</b>	<b>76.2425</b>	498	250
Food	Koza	96.36667	99.5	20.25609	52	116
	$\epsilon = 0.1$	99.43333	104	20.6843	55	<b>129</b>
	$\epsilon = 0.01$	<b>103.933</b>	<b>115</b>	<b>18.6768</b>	<b>65</b>	<b>129</b>
Time	Koza	<b>142.933</b>	<b>138</b>	22.27869	<b>191</b>	<b>107</b>
	$\epsilon = 0.1$	162.3	159.5	<b>15.803</b>	210	139
	$\epsilon = 0.01$	161	150	27.3753	221	129

The obtained results, shown in Table 2, prove that Koza’s GP, while quicker than  $\epsilon$ -GP, trails  $\epsilon$ -GP in both moves and food.  $\epsilon$ -GP obtained the best absolute result by eating 129 out of 156 pieces of food, with 392 moves for  $\epsilon$  value of 0.01 and 410 moves for  $\epsilon$  value of 0.1. With an  $\epsilon$  value of 0.01,  $\epsilon$ -GP was very consistent (low standard deviation) and scored better than the two other sets in both food and moves, although with a slower running time than Koza’s GP. Koza’s GP was not able to collect more than 116 pieces of food, which was achieved with 335 moves in 2 minutes and 28 seconds. As to the fastest run, it was achieved by Koza’s GP, where it collected 52 pieces of food with 493 moves. The lowest number of moves was achieved by  $\epsilon$ -GP with a value of 0.1, where it took 2 minutes and 37 seconds, collecting 55 pieces of food in the process (the lowest score in all runs).

## 7 CONCLUSIONS

The previous section shows promising results, as  $\epsilon$ -GP was shown to simultaneously optimize two objectives, with the algorithm guaranteeing competitive results in all objectives. These results are encouraging for future work as well, as genetic programming, up to our knowledge, has never been used to solve a problem with many (more than 3) objectives using an approximate measure.

Consistency within stochastic algorithms is usually a problem due to the random nature of different runs, but with  $\epsilon$ -GP with an  $\epsilon$  value of 0.01,

this is decreased to an acceptable degree. As problems increase in difficulty, the tolerance of a high  $\epsilon$  value starts to decrease and problems can take longer times to find high-quality solutions and can face a possibility of falling into local minima due to the discarding of many solutions. Therefore, choosing the value of  $\epsilon$  is very important.

This paper serves as an introduction to further work that will test  $\epsilon$ -GP on problems with more than 2 objectives. Furthermore, future work includes the following:

- The value of  $\epsilon$  can be input from the user or dynamically computed.  $\epsilon$  can also be changed to be variable for each objective.
- A more detailed study with better test-set problems that contain more objectives is needed to prove that  $\epsilon$ -GP is an efficient many-objective optimizer;

## REFERENCES

- Abido, M., 2003. A niched Pareto genetic algorithm for multiobjective environmental/economic dispatch. *Electrical Power and Energy Systems*, February, 25(2), pp. 97-105.
- Corne, D. W., Jerram, N. R., Knowles, J. D. & Oates, M. J., 2001. *PESA-II: Region-based Selection in Evolutionary Multiobjective Optimization*. San Francisco, s.n.
- Corne, D. W., Knowles, J. D. & Oates, M. J., 2000. *The Pareto Envelope-based Selection Algorithm for Multiobjective Optimization*. Paris, s.n., pp. 839-848.
- Deb, K., Pratap, A., Agarwal, S. & Meyarivan, T., 2002. A fast and elitist multiobjective genetic algorithm: NSGA-II. *Evolutionary Computation*, April, 6(2), pp. 182-197.
- Fonseca, C. M. & Fleming, P. J., 1993. *Genetic Algorithms for multiobjective optimization: formulation, discussion and generalization*. San Mateo, s.n.
- Khare, V., Yao, X. & Deb, K., 2003. *Performance Scaling of Multi-objective Evolutionary Algorithms*. Faro, s.n., pp. 376-390.
- Knowles, J. & Corne, D., 2000. Approximating the nondominated front using the Pareto Archived Evolution Strategy. *Evolutionary Computation*, 8(2), pp. 149-172.
- Koza, J. R., 1992. *Genetic Programming: On the Programming of Computers by Means of Natural Selection*. 1st ed. London: A Bradford Book.
- Laumanns, M., Thiele, L., Deb, K. & Zitzler, E., 2002. Combining Convergence and Diversity in Evolutionary Multiobjective Optimization. *Evolutionary Computation*, September, 10(3), pp. 263-283.
- Li, M., Yang, S., Liu, X. & Shen, R., 2013. *A Comparative Study on Evolutionary Algorithms for Many-Objective Optimization*. Sheffield, s.n., pp. 261-275.
- Schaffer, J. D., 1985. *Multiple objective optimization with vector evaluated genetic algorithms*. Pittsburgh, s.n., pp. 93-100.
- Srinivas, N. & Deb, K., 1994. Multiobjective optimization using Nondominated sorting in genetic algorithms. *Evolutionary Computation*, September, 2(3), pp. 221-248.
- Zitzler, E., Laumanns, M. & Thiele, L., 2001. *SPEA2: Improving the Strength Pareto Evolutionary Algorithm*, Zurich: s.n.
- Zitzler, E. & Thiele, L., 1999. Multiobjective Evolutionary Algorithms: A comparative Case study and the Strength Pareto Evolutionary Algorithm. *Evolutionary Computation*, November, 3(4), pp. 257 - 271 .