

EVOLUTIONARY OPTIMIZATION OF CONSTRAINED PROBLEMS

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ABSTRACT

In this paper we discuss a construction of Genocop II, a hybrid optimization system for general nonlinear programming problems. We present the first experimental results of the system on five test cases. These include a variety of objective functions with nonlinear constraints. The results are encouraging.

1. Introduction

The general nonlinear programming problem \mathcal{NP} is to find \bar{X} so as to

$$\text{optimize } f(\bar{X}), \bar{X} = (x_1, \dots, x_q) \in R^q,$$

subject to $p \geq 0$ equations:

$$c_i(\bar{X}) = 0, i = 0, \dots, p,$$

and $m - p \geq 0$ inequalities:

$$c_i(\bar{X}) \leq 0, i = p + 1, \dots, m.$$

In this paper we discuss a new hybrid system, Genocop II, to solve optimization problems in this class. The name Genocop is taken from the expression “GEnetic algorithm for Numerical Optimization for COnstrained Problems.” The first version of the Genocop system, Genocop I, handled only linear constraints^{17,18,19}, whereas the current version, Genocop II, handles any set of constraints.

The concept of the presented system is based on the ideas taken from the recent developments in area of optimization³ combined with iterative execution of the Genocop I; these executions are controlled by a temperature of the system (which can be also interpreted as a variable penalty coefficient).

The prototype of the new system Genocop II was run on several test-cases; the results of experiments are presented and compared with known optima.

The paper is organized as follows. The next section summarizes one traditional approach to solve nonlinear programming problems, which was adopted as a basis

for Genocop II. Section 3 introduces the system Genocop II, and the following section discusses several nonlinear test cases together with the experimental results of the Genocop II. The final section contains conclusions and some directions for future work.

2. A Traditional Calculus Based Method

Calculus-based methods assume that the objective function $f(\bar{X})$ and all constraints are twice continuously differentiable functions of \bar{X} . The general approach of most methods is to transform the nonlinear problem \mathcal{NP} into a sequence of solvable subproblems. The amount of work involved in a subproblem varies considerably among methods. These methods require explicit (or implicit) second derivative calculations of the objective (or transformed) function, which in some methods can be ill-conditioned and cause the algorithm to fail.

During the last 30 years there has been considerable research directed toward the nonlinear optimization problems and progress has been made in theory and practice¹². Several approaches have been developed in this area, among these are: the sequential quadratic penalty function^{7,3}, recursive quadratic programming method⁶, penalty trajectory methods²⁰, and the SOLVER method¹⁰. In this section, we discuss briefly one of these approaches, the sequential quadratic penalty function method.

The method replaces a problem \mathcal{NP} by the problem \mathcal{NP}' :

$$\text{optimize } F(\bar{X}, r) = f(\bar{X}) + \frac{1}{2r} \bar{C}^T \bar{C},$$

where $r > 0$ and \bar{C} is a vector of all active constraints c_1, \dots, c_ℓ .

Fiacco and McCormick¹⁵ have shown that the solutions of \mathcal{NP} and \mathcal{NP}' are identical in the limit as $r \rightarrow 0$. It was thought that \mathcal{NP}' could then be solved simply by minimizing $F(\bar{X}, r)$ for a sequence of decreasing positive values of r by Newton's method⁹. This hope, however, was short-lived, because minimizing $F(\bar{X}, r)$ proved to be extremely inefficient for the smaller values of r ; it was shown by Murray²⁰ that this was due to the Hessian matrix of $F(\bar{X}, r)$ becoming increasingly ill-conditioned as $r \rightarrow 0$. As there seemed to be no obvious way of overcoming this problem, the method gradually fell into disuse. More recently, Broyden and Attia^{8,7} offered a method that overcomes the numerical difficulties associated with the simple quadratic penalty function. The computation of the search direction does not require the solution of any system of linear equations, and can thus be expected to require less work than is needed for some other algorithms. The method also provides an automatic technique for calculating the initial value for the parameter r and its successive values⁷.

3. The System Genocop II

The technique discussed in the previous section, together with the existing system Genocop I, was used in construction of a new system, Genocop II. The structure

of the Genocop II is given in Figure 1. We discuss the steps of this algorithm in the remaining part of this section.

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procedure Genocop II
begin
   $t \leftarrow 0$ 
  split the set of constraints  $C$  into
     $C = L \cup N_e \cup N_i$ 
  select a starting point  $\bar{X}_s$ 
  set the set of active constraints,  $A$  to
     $A \leftarrow N_e \cup V$ 
  set temperature  $\tau \leftarrow \tau_0$ 
  while (not termination-condition) do
    begin
       $t \leftarrow t + 1$ 
      execute Genocop I for the function
         $F(\bar{X}, \tau) = f(\bar{X}) + \frac{1}{2\tau} \bar{A}^T \bar{A}$ 
        with linear constraints  $L$ 
        and the starting point  $\bar{X}_s$ 
      save the best individual  $\bar{X}^*$ :
         $\bar{X}_s \leftarrow \bar{X}^*$ 
      update  $A$ :
         $A \leftarrow A - S \cup V$ ,
      decrease temperature  $\tau$ :
         $\tau \leftarrow g(\tau, t)$ 
    end
  end

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Figure 1: The structure of Genocop II.

There are several steps of the algorithm in the first phase of its execution (before it enters the *while* loop). The parameter t (which counts the number of iterations of the algorithm, i.e., the number of times the algorithm Genocop I is applied) is initialized to zero. The set of all constraints C is divided into three subsets: linear constraints L , nonlinear equations N_e and nonlinear inequalities N_i . A starting point \bar{X}_s (which need not be feasible) for the following optimization process is selected (or a user is prompted for it). The set of active constraints A consists initially of elements of N_e and set $V \subseteq N_i$ of violated constraints from N_i . A constraint $c_j \in N_i$ is violated at point \bar{X} iff $c_j(\bar{X}) > \delta$ ($j = p + 1, \dots, m$), where δ is a parameter of the method. Finally, the initial temperature of the system τ is set to τ_0 (a parameter of the method).

In the main loop of the algorithm we apply Genocop I to optimize a modified function

$$F(\bar{X}, \tau) = f(\bar{X}) + \frac{1}{2\tau} \bar{A}^T \bar{A}$$

with linear constraints L . Note that the initial population for Genocop I consists of *pop_size* identical copies (of the initial point for the first iteration and of the best saved point for subsequent ones); several mutation operators introduce diversity in the population at early stages of the process. When Genocop I converges, its best individual \bar{X}^* is saved and used later as the starting point \bar{X}_s for the next iteration. However, the next iteration is executed with a decreased value of the temperature parameter ($\tau \leftarrow g(\tau, t)$) and a new set of active constraints A :

$$A \leftarrow A - S \cup V,$$

where S and V are subsets of N_i satisfied and violated by \bar{X}^* , respectively.

The mechanism of the algorithm is illustrated on the following example. The problem is to

$$\text{minimize } f(\bar{X}) = x_1 \cdot x_2^2,$$

subject to one nonlinear constraint:

$$c_1 : 2 - x_1^2 - x_2^2 \geq 0.$$

The known global solution is $\bar{X}^* = (-0.816497, -1.154701)$, and $f(\bar{X}^*) = -1.088662$. The starting feasible point is $\bar{X}_0 = (-0.99, -0.99)$. After the first iteration of Genocop II (A is empty) the system converged to $\bar{X}_1 = (-1.5, -1.5)$, $f(\bar{X}_1) = -3.375$. The point \bar{X}_1 violates the constraint c_1 , which becomes active. The point \bar{X}_1 is used as the starting point for the second iteration. The second iteration ($\tau = 10^{-1}$, $A = \{c_1\}$) resulted in $\bar{X}_2 = (-0.831595, -1.179690)$, $f(\bar{X}_2) = -1.122678$. The point \bar{X}_2 is used as the starting point for the third iteration. The third iteration ($\tau = 10^{-2}$, $A = \{c_1\}$) resulted in $\bar{X}_3 = (-0.815862, -1.158801)$, $f(\bar{X}_3) = -1.09985$. The sequence of points \bar{X}_t (where $t = 4, 5, \dots$ is the iteration number of the algorithm) approaches the optimum.

4. Test Cases

In order to evaluate the method of Genocop II, a set of test problems has been carefully selected to indicate the performance of the algorithm and to illustrate that it has been successful in practice. The five test cases include quadratic, nonlinear, and discontinuous functions with several nonlinear constraints.

All runs of the system were performed on SUN SPARC station 2. We used the following parameters for Genocop I in all experiments:

pop_size = 70, $k = 28$ (number of parents in each generation), $b = 2$ (coefficient for non-uniform mutation), $a = 0.25$ (parameter of arithmetical crossover), $\delta = 0.01$ (parameter which determines whether or not a constraint is active).

In most cases, the initial temperature τ_0 was set at 1 (i.e., $g(\tau, 0) = 1$); additionally, $g(\tau, t) = 10^{-1} \cdot g(\tau, t - 1)$.

Genocop II was executed ten times for each test case. For most problems, the number of generations necessary for Genocop I to converge was 1000 (more difficult problems required larger number of iterations). We did not report the computational times for these test cases, since we do not have full implementation of Genocop II yet. The actions of the system were simulated by executing its external loop in manual fashion: when Genocop I converges, the best point is incorporated as the starting point for the next iteration, the constraints are checked for their activity status, and the evaluation function is adjusted accordingly.

All test cases and the results of the Genocop II system are reported in the following subsections.

4.1. Test Case #1

The problem¹⁴ is

$$\text{minimize } f(\bar{X}) = 100(x_2 - x_1^2)^2 + (1 - x_1)^2,$$

subject to nonlinear constraints:

$$\begin{aligned} c_1 &: x_1 + x_2^2 \geq 0, \\ c_2 &: x_1^2 + x_2 \geq 0, \end{aligned}$$

and bounds:

$$-0.5 \leq x_1 \leq 0.5, \text{ and } x_2 \leq 1.0.$$

The known global solution is $\bar{X}^* = (0.5, 0.25)$, and $f(\bar{X}^*) = 0.25$. The starting feasible point is $\bar{X}_0 = (0, 0)$.

Genocop II found the exact optimum in one iteration, since none of the nonlinear constraints are active at the optimum.

4.2. Test Case #2

The problem¹¹ is

$$\text{minimize } f(x, y) = -x - y,$$

subject to nonlinear constraints:

$$\begin{aligned} c_1 &: y \leq 2x^4 - 8x^3 + 8x^2 + 2, \\ c_2 &: y \leq 4x^4 - 32x^3 + 88x^2 - 96x + 36, \end{aligned}$$

and bounds:

$$0 \leq x \leq 3 \text{ and } 0 \leq y \leq 4.$$

Iteration number	The best point	Active constraints
0	(0,0)	none
1	(3,4)	c_2
2	(2.06, 3.98)	c_1, c_2
3	(2.3298, 3.1839)	c_1, c_2
4	(2.3295, 3.1790)	c_1, c_2

Table 1: Progress of Genocop II on test case #2; for iteration 0 the best point is the starting point.

The known global solution is $\bar{X}^* = (2.3295, 3.1783)$, and $f(\bar{X}^*) = -5.5079$. The starting feasible point is $\bar{X}_0 = (0, 0)$. The feasible region is almost disconnected.

Genocop II approached the optimum very closely at the 4th iteration. The progress of the system is reported in the table 1.

4.3. Test Case #3

The problem¹¹ is

$$\text{minimize } f(\bar{X}) = (x_1 - 10)^3 + (x_2 - 20)^3,$$

subject to nonlinear constraints:

$$\begin{aligned} c_1 &: (x_1 - 5)^2 + (x_2 - 5)^2 - 100 \geq 0, \\ c_2 &: -(x_1 - 6)^2 - (x_2 - 5)^2 + 82.81 \geq 0, \end{aligned}$$

and bounds:

$$13 \leq x_1 \leq 100 \text{ and } 0 \leq x_2 \leq 100.$$

The known global solution is $\bar{X}^* = (14.095, 0.84296)$, and $f(\bar{X}^*) = -6961.81381$ (see figure 2). The starting point, which is not feasible, is $\bar{X}_0 = (20.1, 5.84)$.

Genocop II approached the optimum very closely at the 12th iteration. The progress of the system is reported in the table 2.

4.4. Test Case #4

The problem⁵ is

$$\text{minimize } f(x_1, x_2) = 0.01x_1^2 + x_2^2,$$

subject to nonlinear constraints:

$$\begin{aligned} c_1 &: x_1x_2 - 25 \geq 0, \\ c_2 &: x_1^2 + x_2^2 - 25 \geq 0, \end{aligned}$$

Figure 2: A feasible space for test case #3.

and bounds:

$$2 \leq x_1 \leq 50 \text{ and } 0 \leq x_2 \leq 50.$$

The global solution is $\bar{X}^* = (\sqrt{250}, \sqrt{2.5}) = (15.811388, 1.581139)$, and $f(\bar{X}^*) = 5.0$. The starting point (not feasible) is $\bar{X}_0 = (2, 2)$.

It is interesting to note that the standard cooling scheme (i.e., $g(\tau, t) = 10^{-1} \cdot g(\tau, t - 1)$) did not produce good results, however, when the cooling process was slowed down (i.e., $g(\tau, 0) = 5$ and $g(\tau, t) = 2^{-1} \cdot g(\tau, t - 1)$), the system approached optimum easily (table 3). This, of course, leads to some questions about how to control the temperature for a given problem: this is one of the topics for future research.

4.5. Test Case #5

The final test problem¹⁴ is

$$\text{minimize } f(\bar{X}) = (x_1 - 2)^2 + (x_2 - 1)^2,$$

subject to a nonlinear constraint:

$$c_1 : -x_1^2 + x_2 \geq 0,$$

and a linear constraint:

$$x_1 + x_2 \leq 2.$$

Iteration number	The best point	Active constraints
0	(20.1, 5.84)	c_1, c_2
1	(13.0, 0.0)	c_1, c_2
2	(13.63, 0.0)	c_1, c_2
3	(13.63, 0.0)	c_1, c_2
4	(13.73, 0.16)	c_1, c_2
5	(13.92, 0.50)	c_1, c_2
6	(14.05, 0.75)	c_1, c_2
7	(14.05, 0.76)	c_1, c_2
8	(14.05, 0.76)	c_1, c_2
9	(14.10, 0.87)	c_1, c_2
10	(14.10, 0.86)	c_1, c_2
11	(14.10, 0.85)	c_1, c_2
12	(14.098, 0.849)	c_1, c_2

Table 2: Progress of Genocop II on test case #3; for iteration 0 the best point is the starting point.

Iteration number	The best point	Active constraints
0	(2,2)	c_1, c_2
1	(3.884181, 3.854748)	c_1
2	(15.805878, 1.581057)	c_1
3	(15.811537, 1.580996)	c_1

Table 3: Progress of Genocop II on test case #4; for iteration 0 the best point is the starting point.

The global solution is $\bar{X}^* = (1, 1)$ and $f(\bar{X}^*) = 1$. The starting (feasible) point is $\bar{X}_0 = (0, 0)$.

Genocop II approached the optimum very closely at the 6th iteration. The progress of the system is reported in the table 4.

5. Conclusions

There are several interesting points connected with the above method. First, as with any other method based on genetic algorithms, it does not require any implicit (or explicit) calculations of gradient or Hessian matrix of the objective function and constraints. Consequently, the method does not suffer from the ill-conditioned Hessian problem usually associated with some calculus-based methods.

Any genetic algorithm can be used in place of Genocop I for the inner loop of

Iteration number	The best point	Active constraints
0	(0, 0)	c_1
1	(1.496072, 0.503928)	c_1
2	(1.020873, 0.979127)	c_1
3	(1.013524, 0.986476)	c_1
4	(1.002243, 0.997757)	c_1
5	(1.000217, 0.999442)	c_1
6	(1.000029, 0.999971)	c_1

Table 4: Progress of Genocop II on test case #5; for iteration 0 the best point is the starting point.

Genocop II. In such a case all constraints (linear and nonlinear) should be considered for placement in the set of active constraints A (the elements of L should be distributed between N_e and N_i). However, such method is much slower and less effective: for efficiency reasons, it is much better to process linear constraints separately (as it is done in Genocop I).

Further research includes experimenting with (1) problems of higher dimensions, (2) different cooling schemes and (3) different values of parameter δ which decides whether a constraint is active or not.

It might be worthwhile to search for other constraint handling paradigms as well. Recently, two interesting new approaches were proposed based on the behavioural memory paradigm and the re-mapping the fitness measures. The first method ²² acts in two stages: (1) a population evolves with some standard GA, where the fitness function is related to the constraint satisfaction, and (2) the final population from the previous stage (viewed as a memory containing some essential information about the constraints) is accepted as initial population for a GA with the objective function as fitness function, which is overridden by assigning zero fitness whenever the constraints are not satisfied. The second method ²¹ re-maps the fitness measures so that all feasible points have higher fitness than infeasible points (which ensures that feasible points are always preferred).

Also, it might be interesting to experiment with “adaptive penalty functions”. After all, probabilities of applied operators might be adaptive (as in evolution strategies ²³); some initial experiments indicate that adaptive population sizes may have some merit ²; so the idea of adaptive penalty functions may deserve some attention. In its simplest version, a penalty coefficient would be part of the solution vector and undergo all operators (as opposed the idea of Genocop II, where such penalty coefficient is changed on regular basis).

Currently, a complete version of GENOCOP II is being implemented, which would include (as options) all above constraint handling methods. Interesting comparisons between these approaches should be reported shortly.

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