

Asymptotic Convergence of Metaheuristics for Multiobjective Optimization Problems

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Abstract

This paper analyzes the convergence of metaheuristics used for multiobjective optimization problems in which the transition probabilities use a uniform mutation rule. We prove that these algorithms converge only if elitism is used.

Key Words: Metaheuristics, multiobjective optimization, multiobjective metaheuristics.

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1 Introduction

This paper concerns metaheuristic algorithms (MhAs) for multiobjective optimization problems (MOPs) (see [1]). For MhAs that use a uniform mutation rule we show that the associated Markov chain converges geometrically to its stationary distribution, but not necessarily to the MOP's optimal solution set. Convergence to the optimal solution set is ensured only if elitism is used.

MhAs are a standard tool to study both single-objective and MOPs. The convergence of a MhA in the single-objective case is reasonably well understood; see [8], for instance. For MOPs, however, the situation is quite different, and to the best of the authors' knowledge, the existing results deal with extremely particular cases; see for example, [9]. This paper is then, the first attempt to deal with the convergence of a general class of MhAs in the context of multiobjective optimization.

The remainder of this paper is organized as follows. Section 2 introduces the MOP we are concerned with. The class of MhAs we are interested on are described in Section 3, together with our main results. These results are proved in Section 4. We conclude in Section 5 with some general remarks and some possible paths of future research.

2 The Multiobjective Optimization Problem

To compare vectors in \mathbb{R}^d we will use the standard Pareto order defined as follows.

If $\vec{u} = (u_1, \dots, u_d)$ and $\vec{v} = (v_1, \dots, v_d)$ are vectors in \mathbb{R}^d , then

$$\vec{u} \preceq \vec{v} \iff u_i \leq v_i \ \forall i \in \{1, \dots, d\}.$$

This relation is a partial order. We also write $\vec{u} \prec \vec{v} \iff \vec{u} \preceq \vec{v}$ and $\vec{u} \neq \vec{v}$.

Definition 1:

Let X be a set and $F : X \longrightarrow \mathbb{R}^d$ a given vector function with components $f_i : X \longrightarrow \mathbb{R}$ for each $i \in \{1, \dots, d\}$. The multiobjective optimization problem (MOP) we are concerned with is to find $x^* \in X$ such that

$$F(x^*) = \min_{x \in X} F(x) = \min_{x \in X} [f_1(x), \dots, f_n(x)], \quad (1)$$

where the minimum is understood in the sense of the Pareto order.

Definition 2:

A point $x^* \in X$ is called a *Pareto optimal solution* for the MOP (1) if there is no $x \in X$ such that $F(x) \prec F(x^*)$. The set

$$\mathcal{P}^* = \{x \in X : x \text{ is a Pareto optimal solution}\}$$

is called the *Pareto optimal set*, and its image under F , i.e.

$$F(\mathcal{P}^*) := \{F(x) : x \in \mathcal{P}^*\},$$

is the *Pareto front*.

As we are concerned with a MhA in which the elements are represented by strings of length l with 0 or 1 in each entry, in the remainder of this paper we will replace X with the *finite* set \mathcal{IB}^l , where $\mathcal{IB} = \{0, 1\}$.

3 Metaheuristic Algorithms

The MhAs are techniques in which there is a population that evolves applying some operations to the current population to obtain the next one. Some of these operations are:

- mutation

- selection
- crossover
- reordering

The MhAs we are interested on are modeled as Markov chains with transition probabilities that use uniform mutation and possibly other operations. This mutation is made with a parameter or probability p_m , which is positive and less than $1/2$, i.e.

$$p_m \in (0, 1/2). \quad (2)$$

In some cases this mutation can be made with two or more parameters, namely the population is divided into subpopulations to each of which a different mutation parameter is applied.

Some examples of this MhAs are

- genetic algorithms (see [6]),
- evolution strategies (see [10]),
- evolutionary programming (see [5, 4]),
- artificial immune systems (see [2, 7]).

The algorithm we are concerned with is modeled as a Markov chain $\{X_k : k \geq 0\}$, whose state space S is the set of all possible populations of n individuals, each one represented by a bit string of length l . Hence $S = \mathcal{B}^{nl}$, where $\mathcal{B} = \{0, 1\}$ and S is the set of all possible vectors of n entries, each of which is a string of length l with 0 or 1 in each entry.

Let $i \in S$ be a state, so that i can be represented as

$$i = (i_1, i_2, \dots, i_n),$$

where each i_s is a string of length l of 0's and 1's.

The chain's transition probability is given by

$$P_{ij} = \mathbb{P}(X_{k+1} = j \mid X_k = i).$$

Thus the transition matrix is of the form

$$P = (P_{ij}) = LM, \tag{3}$$

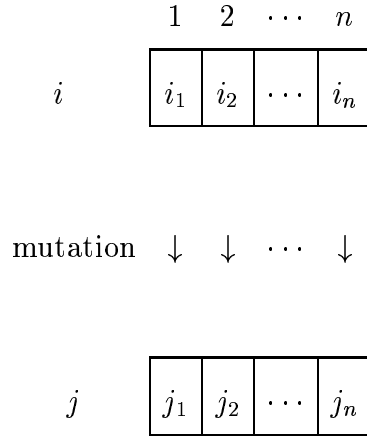
where M is the transition matrix corresponding to the mutation operation and L represents the other operations.

Note that these matrices are stochastic, i.e. $L_{ij} \geq 0$, $M_{ij} \geq 0$ for all i, j , and for each $i \in S$

$$\sum_{s \in S} L_{is} = 1 \quad \text{and} \quad \sum_{s \in S} M_{is} = 1. \quad (4)$$

The Mutation Probability

To calculate the mutation probability from the state i to state j we use that the individual i_s is transformed into the individual j_s applying uniform mutation (i.e., each entry of i_s is transformed into the corresponding one of j_s with probability p_m), as in the following scheme.



Thus, for each individual in the population the mutation probability can

be calculated as

$$p_m^{H(i_s, j_s)} (1 - p_m)^{l - H(i_s, j_s)} \quad \forall s \in \{1, \dots, n\},$$

where $H(i_s, j_s)$ is the Hamming distance between i_s and j_s .

Hence the mutation probability from i to j is:

$$M_{ij} = \prod_{s=1}^n p_m^{H(i_s, j_s)} (1 - p_m)^{l - H(i_s, j_s)} \quad (5)$$

Using Elitism

We say that we are using *elitism* in an algorithm (or a MhA in our case) if we use an extra set, called the *elite* set, in which we put the “best” elements (i.e., the nondominated elements of the state in our case). This elite set usually does not participate in the evolution, since it is used only to store the nondominated elements.

After each transition, we apply an *elitism operation* that accepts a new state if there is an element in the population that improves some element in the elite set.

If we are using elitism, the representation of the states changes to the

following form:

$$\hat{i} = (i^e; i) = (i_1^e, \dots, i_r^e; i_1, \dots, i_n),$$

where i_1^e, \dots, i_r^e are the members of the elite set of the state, r is the number of elements in the elite set and we assume that the cardinality of \mathcal{P}^* is greater than or equal to r . In addition we assume that $r \leq n$.

Note that in general i_1^e, \dots, i_r^e are not necessarily the “best” elements of the state \hat{i} , but after applying the elitism operation in i^e they are the “best” elements of the state.

Let \hat{P} be the transition matrix associated with the new states. If all the elements in the elite set of a state are Pareto optimal, then any state that contains an element in the elite set that is not a Pareto optimal will not be accepted, i.e.

$$\text{if } \{i_1^e, \dots, i_r^e\} \subset \mathcal{P}^* \text{ and } \{j_1^e, \dots, j_r^e\} \not\subset \mathcal{P}^* \text{ then } \hat{P}_{ij} = 0. \quad (6)$$

Main Results

Before stating our main results we introduce the definition of convergence of an algorithm, which uses the following notation: if $V = (v_1, v_2, \dots, v_n)$ is a vector, then $\{V\}$ denotes the set of entries of V , i.e.

$$\{V\} = \{v_1, v_2, \dots, v_n\}.$$

Definition 3:

Let $\{X_k : k \geq 0\}$ be the Markov chain associated to an algorithm. We say that the algorithm converges to \mathcal{P}^* with probability 1 if

$$\mathbb{P}(\{X_k\} \subset \mathcal{P}^*) \rightarrow 1 \text{ as } k \rightarrow \infty.$$

In the case that we are using elitism we replace X_k by X_k^e , the elite set of the state (i.e. if $X_k = i$ then $X_k^e = i^e$)

In the rest of the paper we will assume, for greater generality, that the population of the MhA is divided in two subsets on which we apply different

mutation parameters. In the first subset we apply the parameter p_m and in the second ρ_m . We assume that

$$p_m, \rho_m \in (0, 1/2). \quad (7)$$

Let's assume that the first subset has n_1 individuals, so that the second subset has $n - n_1$ individuals. Thus, for each individual in the first subset of the population the mutation probability can be calculated as

$$p_m^{H(i_s, j_s)} (1 - p_m)^{l - H(i_s, j_s)} \quad \forall s \in \{1, \dots, n_1\},$$

and for the second subset we have

$$\rho_m^{H(i_s, j_s)} (1 - \rho_m)^{l - H(i_s, j_s)} \quad \forall s \in \{n_1 + 1, \dots, n\}.$$

Now, instead of (5) the mutation probability from i to j is:

$$M_{ij} = \prod_{s=1}^{n_1} p_m^{H(i_s, j_s)} (1 - p_m)^{l - H(i_s, j_s)} \prod_{s=n_1+1}^n \rho_m^{H(i_s, j_s)} (1 - \rho_m)^{l - H(i_s, j_s)} \quad (8)$$

Theorem 1:

Let P be the transition matrix of a MhA. Then P has a stationary distribution π such that

$$|P_{ij}^k - \pi_j| \leq (1 - \xi)^{k-1} \quad \forall i, j \in S \quad \forall k = 1, 2, \dots, \quad (9)$$

where $\xi = 2^{nl} p_m^{n_1 l} \rho_m^{(n-n_1)l}$. Moreover, π has all entries positive.

Theorem 1 states that P^k converges geometrically to π . Nevertheless in spite of this result, the convergence of the MhA to the Pareto optimal set cannot be guaranteed. In fact, from Theorem 1 and using the fact that π has all entries positive we will immediately deduce the following.

Corollary 1:

The MhA does not converge.

To ensure convergence of the MhA we need to use elitism.

Theorem 2:

The MhA using elitism converges.

4 Proofs

We first recall some standard definitions and results.

Definition 4:

A stochastic matrix P is said to be *primitive* if there exists $k > 0$ such that the entries of P^k are all positive.

The next result gives an upper bound on the rate of convergence of P^k as $k \rightarrow \infty$. We will use it to show the existence of the stationary distribution in Theorem 1.

Lemma 1:

Let N be the cardinality of S , and let P_{ij}^k be the entry ij of P^k . Suppose that there exists an integer $\nu > 0$ and a set J of $N_1 \geq 1$ values of j such that

$$\min_{\substack{1 \leq i \leq N \\ j \in J}} P_{ij}^\nu = \delta > 0.$$

Then there are a numbers $\pi_1, \pi_2, \dots, \pi_{N_1}$ such that

$$\lim_{k \rightarrow \infty} P_{ij}^k = \pi_j \quad \forall i = 1, \dots, N, \quad \text{with} \quad \pi_j \geq \delta > 0, \quad \forall j \in J,$$

and $\pi_1, \pi_2, \dots, \pi_{N_1}$ form a set of stationary probabilities. Moreover

$$|P_{ij}^k - \pi_j| \leq (1 - N_1\delta)^{\frac{k}{\nu}-1} \quad \forall k = 1, 2, \dots$$

Proof See, for example, [3, p. 173].

The next lemma will allow us to use Lemma 1.

Lemma 2:

Let P be the transition matrix of the MhA. Then

$$\min_{i,j \in S} P_{ij} = p_m^{n_1 l} \rho_m^{(n-n_1)l} > 0 \quad \forall i, j \in S, \quad (10)$$

and therefore P is primitive.

Proof

By (7) we have

$$p_m < \frac{1}{2} < 1 - p_m, \quad \rho_m < \frac{1}{2} < 1 - \rho_m.$$

Thus, from (8),

$$\begin{aligned}
M_{ij} &= \prod_{s=1}^{n_1} p_m^{H(i_s, j_s)} (1 - p_m)^{l - H(i_s, j_s)} \prod_{s=n_1+1}^n \rho_m^{H(i_s, j_s)} (1 - \rho_m)^{l - H(i_s, j_s)} \\
&> \prod_{s=1}^{n_1} p_m^l \prod_{s=n_1+1}^n \rho_m^l \\
&= p_m^{n_1 l} \rho_m^{(n - n_1) l}
\end{aligned}$$

On the other hand, by (3) and (4)

$$\begin{aligned}
P_{ij} &= \sum_{s \in S} R_{is} M_{sj} \\
&\geq p_m^{n_1 l} \rho_m^{(n - n_1) l} \sum_{s \in S} R_{is} \\
&= p_m^{n_1 l} \rho_m^{(n - n_1) l} > 0,
\end{aligned}$$

To verify (10), observe that P_{ij} attains the minimum in (10) if i has 0 in all entries and j has 1 in all entries. Thus the desired conclusion follows. \blacksquare

Proof of Theorem 1

From Lemma 2, P is primitive. Moreover, because (10) holds for all $j \in S$ we have that $N_1 = N = 2^{nl}$ and $\nu = 1$. Thus, by Lemma 1, P has a stationary

distribution π with all entries positive and we get (9). ■

Before proving Theorem 2 we give some definitions and preliminary results.

Definition 5:

Let X be as in Definition 1. We say that X is *complete* if for each $x \in X \setminus \mathcal{P}^*$ there exists $x^* \in \mathcal{P}^*$ such that $F(x^*) \preceq F(x)$. For instance, if X is finite then X is complete.

Let $i, j \in S$ be two arbitrary states, we say that i *leads* to j , and write $i \rightarrow j$, if there exists an integer $k \geq 1$ such that $P_{ij}^k > 0$. If i does not lead to j we write $i \not\rightarrow j$.

We call a state i *inessential* if there exists a state j such that $i \rightarrow j$ but $j \not\rightarrow i$. Otherwise, the state i is called *essential*.

We denote the set of essential states by E and the set of inessential states by I . Clearly,

$$S = E \cup I.$$

We say that P is in *canonical form* if it can be written as

$$P = \begin{pmatrix} P_1 & 0 \\ R & Q \end{pmatrix}.$$

Observe that P can put in this form by reordering the states, that is, the essential states at the beginning and the inessential states at the end. In this case, P_1 is the matrix associated with the transitions between essential states, R with transitions from inessential to essential states, and Q with transitions between inessential states.

Note that P^k has a Q^k in the position of Q in P , i.e.

$$P^k = \begin{pmatrix} P_1^k & 0 \\ R_k & Q^k \end{pmatrix},$$

where R_k is a matrix that depends of P_1 , Q and R .

Now we present some results that will be essential in the proof of Theorem 2.

Lemma 3:

Let P be a stochastic matrix, and let Q be the submatrix of P associated

with transitions between inessential states. Then, as $k \rightarrow \infty$,

$$Q^k \rightarrow 0 \text{ elementwise geometrically fast.}$$

Proof See, for instance, [11, p.120]. ■

As a consequence of Lemma 3 we have the following.

Corollary 2:

For any initial distribution,

$$\mathbb{P}(X_k \in I) \rightarrow 0 \text{ as } k \rightarrow \infty.$$

Proof

For any initial distribution vector p_0 , let $p_0(I)$ be the subvector that corresponds to the inessential states. Then, by Lemma 3,

$$\mathbb{P}(X_k \in I) = p_0(I)' Q^k \mathbf{1} \rightarrow 0 \text{ as } k \rightarrow \infty.$$

■

Proof of Theorem 2

By Corollary 2, it suffices to show that the states that contain elements in the elite set that are not Pareto optimal are inessential states. To this end, first note that $X = \mathcal{B}^l$ is complete, because it is finite.

Now suppose that there is a state $\hat{i} = (i^e; i)$ in which the elite set contains elements $i_{s_1}^e, \dots, i_{s_k}^e$ that are not Pareto optimal. Then, as X is complete, there are elements, say $j_{s_1}^e, \dots, j_{s_k}^e \in \mathcal{P}^*$, that dominate $i_{s_1}^e, \dots, i_{s_k}^e$, respectively.

Take $\hat{j} = (j^e; j)$ such that all Pareto optimal points of i^e are in j^e and replace the other elements of i^e with the corresponding $j_{s_1}^e, \dots, j_{s_k}^e$. Thus all the elements in j^e are Pareto optimal.

Now let

$$j = (j_1^e, \dots, j_r^e, \underbrace{i_{s_1}^e, \dots, i_{s_1}^e}_{n-r \text{ copies}}).$$

By Lemma 2 we have $i \rightarrow j$. Hence with positive probability we can pass from (i^e, i) to (i^e, j) , and then we apply the elitism operation to pass from (i^e, j) to (j^e, j) . This implies that $\hat{i} \rightarrow \hat{j}$. On the other hand, using (6), $\hat{j} \not\rightarrow \hat{i}$ and therefore \hat{i} is an inessential state.

Finally, from Corollary 2 we have

$$\mathbb{P}(\{X_k^e\} \subset \mathcal{P}^*) = \mathbb{P}(X_k \in E) = 1 - \mathbb{P}(X_k \in I) \rightarrow 1 - 0 = 1$$

as $k \rightarrow \infty$. This completes the proof of Theorem 2. ■

5 Conclusions and Future Work

We have presented a general convergence analysis of a MhA for MOPs in which uniform mutation is used. It was proven in Theorem 2 that it is necessary to use elitism to ensure that our algorithm converges. This result is of course reassuring, but it is not quite complete in the sense that we have been unable to provide a result such as in (9), on the speed of convergence. The latter fact as well as a convergence analysis of a MhA with nonuniform mutation rule, require further research.

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