EVOLUTIONARY MULTI-OBJECTIVE SCHEDULING PROCEDURES IN NON-STANDARDIZED PRODUCTION PROCESSES

PROCEDIMIENTOS DE PROGRAMACIÓN EVOLUTIVA MULTI-OBJETIVO EN PROCESOS PRODUCTIVOS NO ESTANDARIZADOS

MARIANO FRUTOS

Doctor en Ingeniería, Universidad Nacional del Sur y CONICET, Argentina, mfrutos@uns.edu.ar

FERNANDO TOHMÉ

Doctor en Economía, Universidad Nacional del Sur y CONICET, Argentina, ftohme@criba.edu.ar

ABSTRACT: Scheduling problems can be seen as multi-objective optimization problems (MOPs), involving the simultaneous satisfaction of several goals related to the optimal design, coordination, and management of tasks. The complexity of the goal functions and of the combinatorial methods used to find analytical solutions to them is quite high. The search for solutions (Pareto-optima) is better served by the use of genetic algorithms (GAs). In this paper, we analyze the performance of the non-dominated sorting genetic algorithm II (NSGAII), strength Pareto evolutionary algorithm II (SPEAII), and their predecessors, NSGA and SPEA, when these are devoted to scheduling tasks in non-standardized production activities.

KEYWORDS: Job-shop scheduling, multi-objective optimization, Pareto frontier, memetic algorithm, local search

RESUMEN: En los problemas de programación de la producción que involucran diseñar, coordinar, administrar y controlar todas las operaciones presentes en el proceso productivo, aparecen numerosos problemas de optimización multi-objetivo (MOPs). Los MOPs constan de varias funciones que suelen ser complejas y evaluarlas puede ser muy costoso. La optimización multi-objetivo es la disciplina que trata de encontrar las soluciones, denominadas Pareto óptimas, a este tipo de problemas. La compleja resolución de los MOPs es debida a las dimensiones del problema, al carácter combinatorio de los algoritmos y a la naturaleza de los objetivos los cuales están vinculados a la eficiencia del sistema. En las últimas décadas muchos MOPs vinculados a la producción han sido tratados con éxito con técnicas de resolución basadas en algoritmos genéticos (GAs). En este trabajo se evalúa a NSGAII (Non-dominated Sorting Genetic Algorithm II), SPEAII (Strength Pareto Evolutionary algorithm II) y a sus antecesores, NSGA y SPEA, en el proceso de planificación de la producción no estandarizada.

PALABRAS CLAVE: Programación job-shop, optimización multi-objetivo, frontera de Pareto, algoritmo memético, búsqueda local

1. INTRODUCTION

The scheduling of job-shop (i.e., non-standardized) production activities requires for one to assign in the best possible way the resources used in those processes [1–4]. This, in turn, demands efficient procedures to optimize decisions in those contexts [5,6]. This job-shop scheduling problem (JSSP) has been classified as NP-Hard, meaning that no polynomial algorithm has been found for solving it. Worse yet, the time required to find a solution grows exponentially with the size of the problem [7,8]. Different alternative presentations of the problem have been advanced, in order to accelerate the search for solutions [9–12]. A common feature of most JSSPs is the presence of at least two conflicting goals that have to be simultaneously optimized [13]. Such multi-objective optimization problems usually have many different solutions.

If we assume that, without loss of generality, all the objectives have to be minimized, a multi-objective optimization problem (MOP) requires finding a vector $\vec{x}^* = [x_1^*,...,x_n^*]^T$ satisfying q inequality constraints $g_i(\vec{x}) \, ^3 \, 0, \, i = 1,...,q$ and p equality constraints $h_i(\vec{x}) = 0, \, i = 1,...,p$ that minimizes $\vec{f}(\vec{x}) = [f_1(\vec{x}),...,f_k(\vec{x})]^T$, where the vector of decision variables is $\vec{x} = [x_1,...,x_n]^T$.

The class of values that satisfy the constraints defines a region of feasible solutions, denoted Ω. That is, any

$\vec{x} \in \Omega$ is a possible solution to the MOP. Then, $\vec{x}^* \in \Omega$ is Pareto optimal if there is no other feasible $\vec{x} \in \Omega$ that improves on any goal without worsening another one. Feasible solutions can be partially ordered in terms of Pareto dominance: a vector $\vec{u} = [u_1,...,u_n]^T$ dominates another vector $\vec{v} = [v_1,...,v_n]^T$ (denoted $\vec{u} \prec \vec{v}$) if and only if $\forall i \in \{1,...,k\}$, $u_i \leq v_i \wedge \exists i \in \{1,...,k\} : u_i < v_i$.

This allows us to define, for a MOP with objective function $\vec{f}(\vec{x})$, the optimal Pareto set $P^* = \{\vec{x} \in \Omega \mid \neg \exists \vec{x}' \in \Omega, \vec{f}(\vec{x}') \prec \vec{f}(\vec{x})\}$. Finally, this set leads to the definition of the Pareto frontier of the problem: $FP^* = \{\vec{f}(\vec{x}), \vec{x} \hat{\imath} P^*\}$. The main goal in the analysis of MOPs is to find the corresponding Pareto frontier. Since it can contain a large number of points, a good solution consists only of a few points, as close as possible from the exact Pareto frontier and uniformly distributed along its contour.

Quite useful tools for the search of this approximate Pareto frontier are evolutionary algorithms [14,15]. In particular, genetic algorithms (GAs) are easy to program and implement [16]. Nevertheless, the high rate of convergence of GAs is costly, since this induces the loss of diversity in the solutions, which is reflected in poorly distributed Pareto frontiers. But if a GA is complemented by an efficient local search method, it yields a procedure that solves multi-objective problems requiring a low number of evaluations of the fitness function. This kind of combined algorithm is called a multi-objective memetic algorithm [17].

2. A MULTI-OBJECTIVE MEMETIC ALGORITHM

We introduce the reader here to a multi-objective memetic algorithm that operates on chromosomes representing the sequence of operations to carry out on different machines, one chromosome for each machine. Each chromosome is coded as an ordered list of integers, representing the order in which the jobs will be performed. That is, with values between 0 and $n!-1$ (n is the total number of jobs to be done), we represent the sequence of jobs in a given machine. For $m = 3$ (m is the number of machines) and $n = 3$, $0 \rightarrow 123$, $1 \rightarrow 132$, $2 \rightarrow 213$, $3 \rightarrow 231$, $4 \rightarrow 312$ and $5 \rightarrow 321$. The initial population consists of individuals in which the chromosomes have randomly assigned genes. The

algorithm starts by decoding and evaluating these individuals. Two functions are evaluated: makespan (the time required to finish all the jobs) [18] (Eq. 1) and tardiness (the longest delay in finishing a job) [19] (Eq. 2).

$$O_1 \rightarrow C_{max} = max\ (C_j) \qquad (1)$$

$$O_2 \rightarrow T_{max} = max\ (T_j) \qquad (2)$$

Here C_j is the date in which job j is finished, while T_j is the delay of job j with respect to its intended finishing date. T_j is computed up from Eq. 3, in which d_j is the due date of delivery.

$$T_j = max\{(C_j - d_j),0\} \qquad (3)$$

in turn, d_j is determined according to Eq. 4, where τ_j^i is operation i's processing time for the j job. Here we take β to be 0.20.

$$d_j = (1+\beta)\sum_{i=1}^{m} \tau_j^i \qquad (4)$$

The evaluation determines the criticality of each machine [20]. According to the evaluation, basic genetic operations are applied on the chromosomes. After that, an improvement operator is applied on the individuals, according to the meta-heuristic method known as simulated annealing (SA) [21]. This technique searches locally for better solutions. SA allows for one to search in less favorable areas of the state space, according to the density of potential solutions found. After this step, the resulting individuals are pooled together and subject to non-dominated sorting genetic algorithm II (NSGAII) [22]. NSGAII applies an elitist strategy together with an explicit mechanism to ensure diversity. The elitist procedure consists of choosing the better individuals from the union of the parent and children populations. Fig. 1 describes the operation of the multi-objective memetic algorithm.

3. COMPARING NSGAII TO OTHER ALGORITHMS

We consider three multi-objective evolutionary algorithms (MOEAs) and compare them to NSGAII. The MOEAs are: non-dominated sorting genetic algorithm (NSGA) [23], strength Pareto evolutionary

algorithm (SPEA) [24], and strength Pareto evolutionary algorithm II (SPEAII) [25].

Generate the initial population (P_0) of size N

Decode and evaluate $O_1(x)$ and $O_2(x)$ for each $x \in P_0$

Assign values r_i and d_i to each $x \in P_0$

Select parents from P_0

Q_0 = Crossover (P_0)

Q'_0 = Mutate (Q_0)

Q''_0 = Search locally (Q'_0)

for i = 0 to (G − 1) **do**

Decode and evaluate $O_1(x)$ and $O_2(x)$ for each $x \in Q''_i$

Assign values r_i and d_i to each $x \in Q''_i$

Select the N best individuals from $P_i \cup Q''_i$

Create the next generation P_{i+1}

Select parents from P_{i+1}

Q_{i+1} = Crossover (P_{i+1})

Q'_{i+1} = Mutate (Q_{i+1})

Q''_{i+1} = Search locally (Q'_{i+1})

end for

end

Figure 1. Multi-objective memetic algorithm

NSGA classifies individuals by layers of undominated individuals, giving each one in a layer the same value of fitness. This value is proportional to the entire population considered. Once obtained a top layer, it is dismissed and a new layer is considered. The process continues until all the population is classified. Since the individuals in the top layer have the highest degree of fitness, they get more attention than the rest of the population. SPEA creates a file containing the undominated solutions found previously. This file is updated, including new undominated solutions, and deleting the solutions that become dominated. For each one a strength value is computed, proportional to the number of solutions dominated by the individual. The fitness of each member is obtained by computing

the strengths of those which dominate it. SPEAII improves over its predecessor by assigning to each possible solution a fitness value that depends both on the number of individuals dominated and the number of those dominating it. It uses also the "closest neighbor" method to assess the density of solutions and guiding the search more efficiently.

4. EXPERIMENTS AND RESULTS

The algorithms discussed above were implemented in a platform and programming language-independent interface for search algorithms (PISA) [26]. PISA is an interface that allows for one to program search algorithms, splitting them in two modules: the variator and the selector. The variator module captures all the specifics of the optimization problem, and its goal is to decode and evaluate the fitness of individuals. The selector module, in turn, is independent of the details of the problem and runs the selection process. Both modules communicate through text files that allow for one to implement the algorithms independently of the underlying programming language and the operations system.

The experiments were run adding a local search stage to each of the aforementioned algorithms. An exploratory analysis showed that the process of improvement of solutions tends to stabilize around the 200th generation. This allowed us to restrict the experiments to stop at the 500th generation, leaving ample room for a late improvement. The experiments were run on a 3.00 GHZ CPU with 1.00 GB RAM. The parameters imposed on the experiments were as follows: the size of the population was 200; the number of generations, 500; the crossover type, uniform; the probability of a crossover, 0.90; the mutation type, two-swap; the probability of mutation, 0.01; the local search type, simulated annealing and; finally, the probability of local search was 0.01. We present the results for problems la01, la02, la03, la04, la05, la06, la07, la08, la09, and la10 [27] in which the goal is to minimize makespan and delay. The procedure is the usual one in these cases [28]. Each algorithm was run 10 times. For each of them, ten classes of undominated solutions were obtained: $P_1, P_2, \ldots, P_{10}$. A super-population $P_T = P_1 \cup P_2 \cup \ldots \cup P_{10}$ was created for each algorithm. From each super-population the undominated solutions were extracted to form the Pareto frontiers Y_{NSGA}, Y_{SPEA}, Y_{SPEAII}, and Y_{NSGAII}. The outcomes can be seen by observing Figs. 2 to 11.
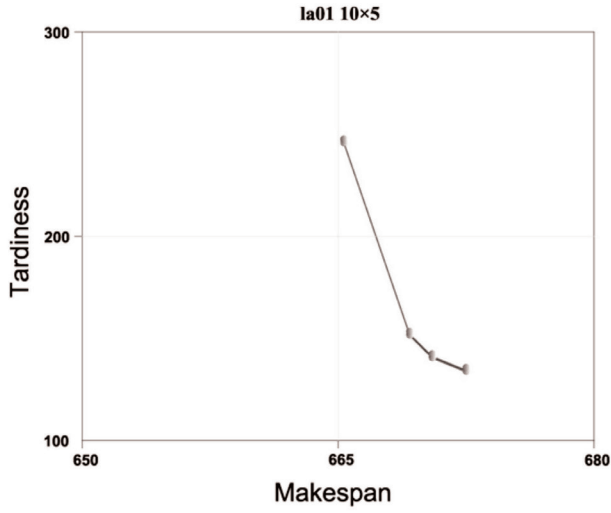
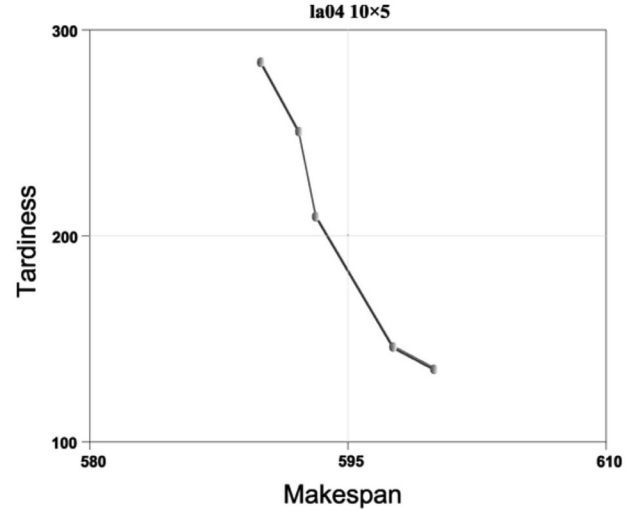Figure 2. la01, Y_{NSGA}(•), Y_{SPEA}(·), Y_{SPEAII}(•) and Y_{NSGAII}(•)

Figure 5. la04, Y_{NSGA}(•), Y_{SPEA}(·), Y_{SPEAII}(•) and Y_{NSGAII}(•)
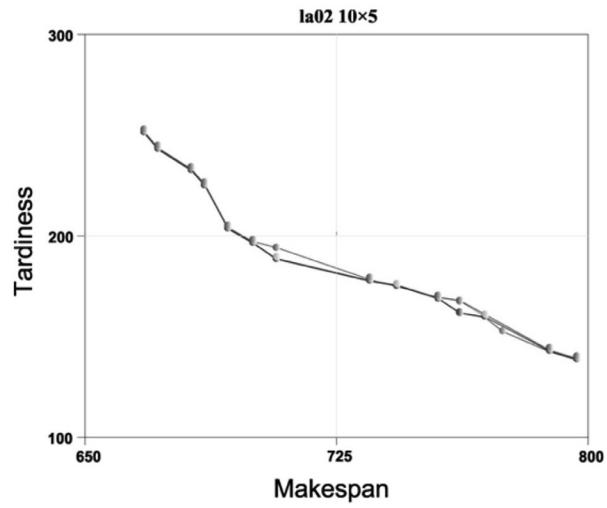
Figure 3. la02, Y_{NSGA}(•), Y_{SPEA}(·), Y_{SPEAII}(•) and Y_{NSGAII}(•)
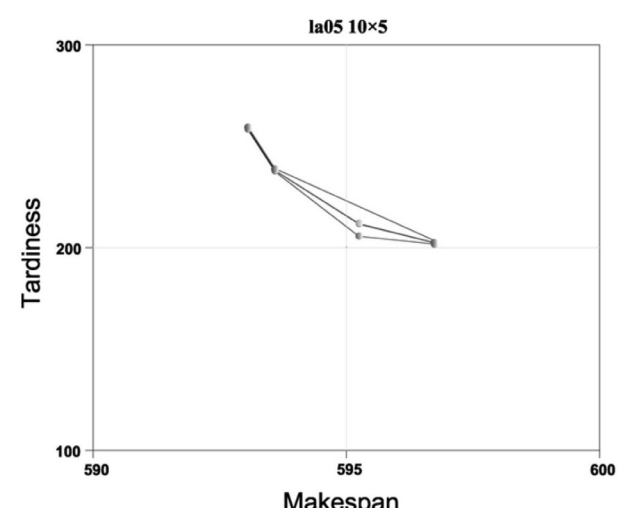
Figure 6. la05, Y_{NSGA}(•), Y_{SPEA}(·), Y_{SPEAII}(•) and Y_{NSGAII}(•)
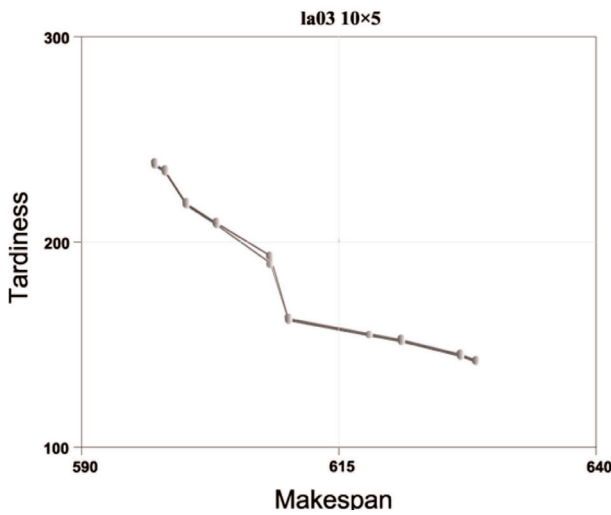
Figure 4. la03, Y_{NSGA}(•), Y_{SPEA}(·), Y_{SPEAII}(•) and Y_{NSGAII}(•)
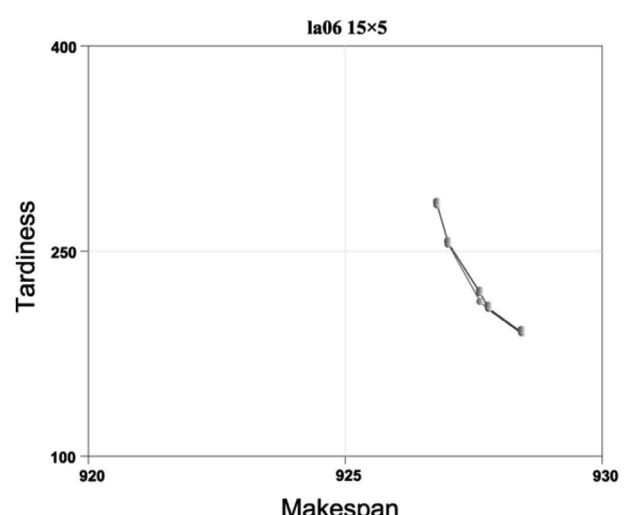
Figure 7. la06, Y_{NSGA}(•), Y_{SPEA}(·), Y_{SPEAII}(•) and Y_{NSGAII}(•)
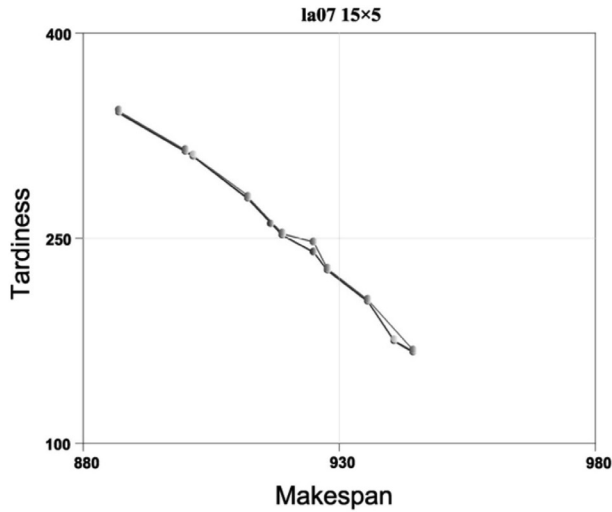
Figure 8. la07, $Y_{NSGA}^{(\bullet)}$, $Y_{SPEA}^{(\cdot)}$, $Y_{SPEAII}^{(\bullet)}$ and $Y_{NSGAII}^{(\bullet)}$
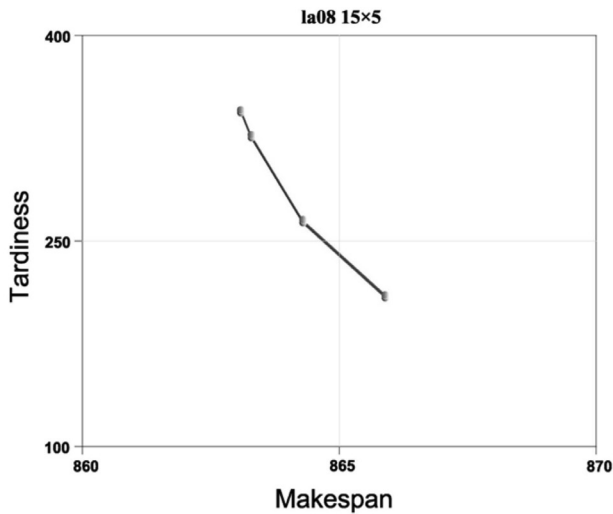
Figure 9. la08, $Y_{NSGA}^{(\bullet)}$, $Y_{SPEA}^{(\cdot)}$, $Y_{SPEAII}^{(\bullet)}$ and $Y_{NSGAII}^{(\bullet)}$
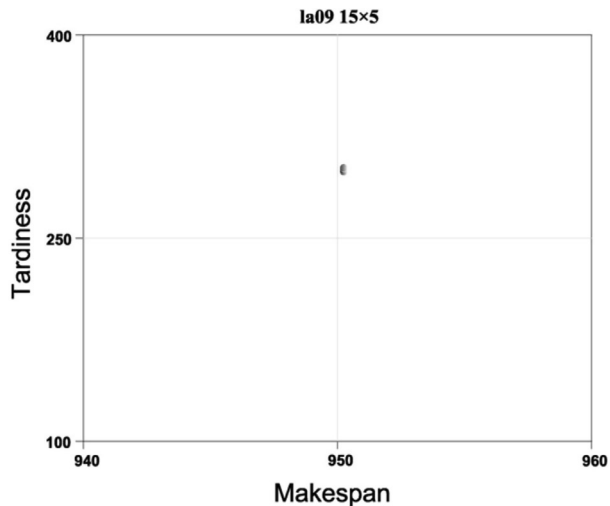
Figure 10. la09, $Y_{NSGA}^{(\bullet)}$, $Y_{SPEA}^{(\cdot)}$, $Y_{SPEAII}^{(\bullet)}$ and $Y_{NSGAII}^{(\bullet)}$
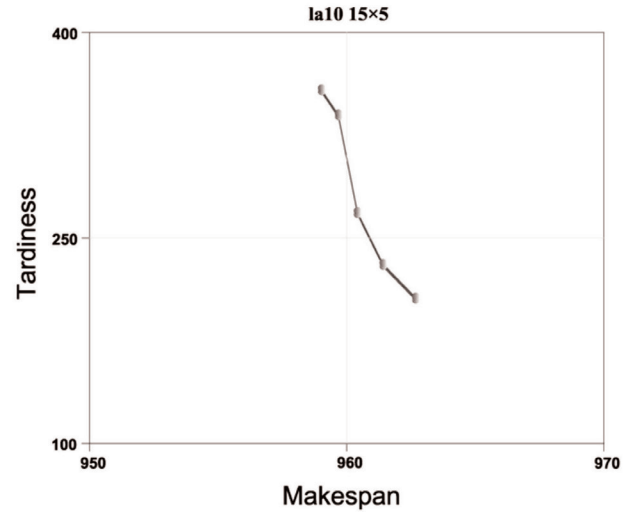
Figure 11. la10, $Y_{NSGA}^{(\bullet)}$, $Y_{SPEA}^{(\cdot)}$, $Y_{SPEAII}^{(\bullet)}$ and $Y_{NSGAII}^{(\bullet)}$

To approximate the optimal Pareto frontier we take $Y_{BK} = Y_{NSGA} \cup Y_{SPEA} \cup Y_{SPEAII} \cup Y_{NSGAII}$ (Y_{BK} for *best known* solutions) and eliminate the dominated solutions. The comparison takes into account which solutions of each algorithm belong to Y_{BK} and the solutions from one algorithm dominating the solutions of another: A solution is effective if it belongs to Y_{BK} and is not dominated by another one in Y_{BK}.

5. COMPARISON PROCEDURE

The algorithms were compared in terms of the resulting values of makespan and delay. For problems la01 (Fig. 2), la04 (Fig. 5), la08 (Fig. 9), la09 (Fig. 10), and la10 (Fig. 11), it can be seen that NSGA, SPEA, NSGAII, and SPEAII have all their solutions in Y_{BK}. In these cases, no algorithm improves over the others. For la02 (Fig. 3) 100% of the solutions of NSGAII are in Y_{BK}. Even if SPEAII is 100% effective, not all of its solutions are in Y_{BK}. SPEA reaches 92.8% of its effectiveness, since from 14 solutions only 1 of them is dominated by an element of Y_{BK}. We can see that NSGAII and SPEAII dominate NSGA in 2 solutions and SPEA in 1. Besides, SPEA dominates NSGA in 1 solution. In la03 (Fig. 4), 100% of the solutions of NSGAII are in Y_{BK}. Despite the 100% effectiveness of SPEAII, not every alternative of Y_{BK} obtains one of its solutions. NSGAII and SPEAII dominate NSGA and SPEA in 1 solution. For la05 (Fig. 6), it can be seen that NSGAII has all its solutions in Y_{BK}. Even if NSGA has 100% of effectiveness, not all of its solutions are in Y_{BK}. NSGAII dominates SPEA and SPEAII in 1 solution.

For la06 (Fig. 7), we have that only NSGAII has 100% of its solutions in Y_{BK}. NSGAII dominates NSGA, SPEA and SPEAII in 1 solution. In the case of la07 (Fig. 8), we see that all of the solutions of SPEAII belong to Y_{BK}. Although NSGAII has 100% effectiveness, not all of its solutions are in Y_{BK}. NSGAII and SPEAII dominate NSGA and SPEA in 1 solution. We can see that NSGAII reaches, in most cases, all the solutions in Y_{BK}. SPEAII also exhibits a good performance, although not a performance as good as NSGAII. NSGA and SPEA, instead, reach less solutions for Y_{BK}. In summary, NSGAII seems to be a better alternative, and justifies its selection as the core GA in our memetic algorithm.

6. CONCLUSIONS

We presented a multi-objective memetic algorithm to solve job-shop scheduling problems (JSSPs) based on NSGAII and simulated annealing. To assess how well this algorithm behaves on this class of NP-hard problems, we run it on a family of well-known JSSPs, arising in non-standardized production contexts. Furthermore, we consider alternative memetic algorithms in which NSGAII is replaced by other multi-objective evolutionary algorithms (MOEAs). In most of the problems, the quality of the solutions found by NSGAII is equal or higher than the results of SPEAII. On the other hand, NSGAII definitely improves over NSGA and SPEA. We can conclude that the multi-objective memetic algorithm here is a good approach for solving JSSPs. Future work involves running this algorithm over other kinds of problems.

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