

A HYBRID TECHNIQUE FOR HANDLING THE FLEXIBLE JOB-SHOP SCHEDULING PROBLEM

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I. INTRODUCTION

To schedule production in a Job-Shop environment means to allocate adequately the available resources. To do that it is necessary to rely on efficient optimization procedures. In fact, the Job-Shop Scheduling Problem is a NP-Hard problem [Ullman, 1975], so ad-hoc algorithms have to be applied to its solution [Frutos et al., 2008]. This is similar to other combinatorial programming problems [Larrazábal, 2003], [Olivera et al., 2006], [Cortés et al., 2004].

Most instances of the Job-Shop Scheduling Problem involve the simultaneous optimization of two usually conflicting goals. This one, like most multi-objective problems, tends to have many solutions. The Pareto frontier reached by an optimization procedure has to contain a uniformly distributed number of solutions close to the ones in the true Pareto frontier. This feature facilitates the task of the expert who interprets the solutions [Kacem et al., 2002].

In this paper we present a Genetic Algorithm linked to a Simulated Annealing procedure able to schedule the production in a Job-Shop manufacturing system [Cortés et al., 2004], [Tsai and Lin, 2003], [Wu et al., 2004].

II. FLEXIBLE JOB-SHOP SCHEDULING PROBLEM

The Job-Shop Scheduling Problem (JSSP) can be described as that of organizing the execution of n jobs on m machines. We assume a finite number of tasks, $\{J_j\}_{j=1}^n$. These tasks must be processed by a finite number of machines $\{M_k\}_{k=1}^m$. To process a task J_j in a machine M_k is denoted by O_{jk}^i , where i indicates the order in which a class of operations $\{S_j\}_{j=1}^n$ is applied on a task J_j . O_{jk}^i requires the uninterrupted use of a machine M_k for a period τ_{jk}^i (the processing time) at a cost u_{jk}^i . In the particular case of Flexible JSSP (FLEX-JSSP), the allocation of O_{jk}^i on M_k is undifferentiated, which means that each O_{jk}^i can be processed by any of the machines in $\{M_k\}_{k=1}^m$. After allocating the operations, we obtain a finite class E of groupings of the O_{jk}^i s on the same machine. We denote each of these groupings as E_k , for $k=1, \dots, m$. A key issue here is the scheduling of activities, i.e. the determination of the starting time t_{jk}^i of each O_{jk}^i .

The FLEX-JSSP demands a procedure to handle its two sub-problems: the allocation of the O_{jk}^i s on the different M_k s and their sequencing, guided by the goals to reach. That is, to find optimal levels of Processing Time (PT*) (see EQ. 1) and Operation Costs (OC*) (see EQ. 2).

$$PT^* \rightarrow \text{Min} : \max (t_{jk}^i + \tau_{jk}^i) \quad (\text{EQ. 1})$$

$$OC^* \rightarrow \text{Min} : \sum_j \sum_i \sum_k x_{jk}^i u_{jk}^i \quad (\text{EQ. 2})$$

where $x_{jk}^i = 1$ if $O_{jk}^i \in E_k$ and 0 otherwise. On the other hand $\sum_k x_{jk}^i = 1$. Besides, $t_{jk}^i = \max (t_{jh}^{(i-1)} + \tau_{jh}^{(i-1)}, t_{pk}^s + \tau_{pk}^s, 0)$ for each pair $O_{jh}^{i-1}, O_{pk}^s \in E_k$ and all machines M_k, M_h and operations S_i, S_s .

III. HYBRID GENETIC ALGORITHM

Due to its many advantages, evolutionary algorithms have become very popular for solving multi-objective optimization problems [Zitzler et al., 2001], [Coello Coello et al., 2002]. Among the evolutionary algorithms used, some of the most interesting are Genetic Algorithms (GA) [Goldberg, 1989].

To represent the individuals, we use a variant of [Wu et al., 2004]. Since the FLEX-JSSP has two subproblems, the Hybrid Genetic Algorithm (HGA) presented here operates over two chromosomes. The first one represents the allocation A_{jk}^i of each O_{jk}^i to every M_k . We denote with values between 0 and $(k-1)$ the allocation of each M_k , that is, for $k=4$, we might have something like $0 \rightarrow M_1, 1 \rightarrow M_2, 2 \rightarrow M_3$ and $3 \rightarrow M_4$ (see TABLE 1A). The second chromosome represents the sequencing of the O_{jk}^i already assigned to each of the M_k ($\forall O_{jk}^i \in E_k$). We denote with values between 0 and $(n!-1)$ the sequence of J_j in each M_k . That is, for $n=3$, we may have $0 \rightarrow J_1 J_2 J_3, 1 \rightarrow J_1 J_3 J_2, 2 \rightarrow J_2 J_1 J_3, 3 \rightarrow J_2 J_3 J_1, 4 \rightarrow J_3 J_1 J_2, 5 \rightarrow J_3 J_2 J_1$ (see TABLE 1B).

| Allocations | | | | Sequences | | | |
|-------------|------------|------------|-----|-----------|---------------|---------------|---|
| J_j | O_{jk}^i | A_{jk}^i | Chr | M_k | Order | Chr | |
| J_1 | O_{1k}^1 | M_4 | 3 | M_1 | $J_2 J_3 J_1$ | 3 | |
| | O_{1k}^2 | M_4 | 3 | | M_2 | $J_2 J_3 J_1$ | 3 |
| | O_{1k}^3 | M_1 | 0 | | M_3 | $J_1 J_2 J_3$ | 0 |
| J_2 | O_{2k}^1 | M_2 | 1 | M_4 | $J_3 J_2 J_1$ | 5 | |
| | O_{2k}^2 | M_2 | 1 | | | | |
| | O_{2k}^3 | M_4 | 3 | | | | |
| J_3 | O_{3k}^1 | M_3 | 2 | | | | |
| | O_{3k}^2 | M_4 | 3 | | | | |
| (A) | | | | (B) | | | |

TABLE I CHROMOSOMES OF ALLOCATIONS AND THE SEQUENCES

The algorithm NSGA-II (Non-Dominated Sorting Genetic Algorithm II) [Deb et al., 2002], creates an initial population, be it random or otherwise. NSGA-II uses an elitist strategy joint with an explicit diversity mechanism. Each individual candidate solution i is assumed to have an associated rank of non-dominance r_i and a distance d_i which indicates the radius of the area in the search space around i not occupied by another solution. A solution i is preferred over j if $r_i < r_j$. When i and j have the same rank, i is preferred if $d_i > d_j$. Let Y_i be an ordered class of individuals with same rank as i and f_j^{i+1} the value for objective j for the individual after i , while f_j^{i-1} is the value

for the individual before i . f_j^{\max} is the maximal value for j among Y_i while f_j^{\min} is the minimal value among Y_i (see FIG. 1). The distances consider all the objective functions and attach an infinite value to the extreme solutions in Y_i . Since these yield the best values for one of the objective functions on the frontier, the resulting distance is the sum of the distances for the objective functions.

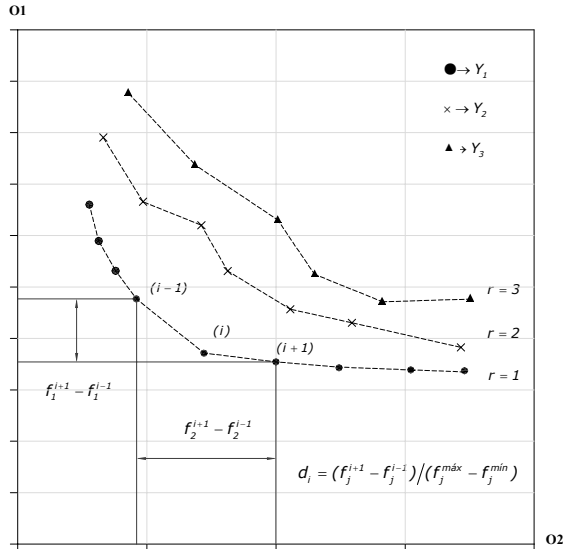


FIG. 1 NON-DOMINATED SORTING GENETIC ALGORITHM II (NON-DOMINANCE RANK R_i AND DISTANCE D_i)

Starting with a population P_t a new population of descendents Q_t obtains. These two populations mix to yield a new one, R_t of size $2N$ (N is the original size of P_t). The individuals in R_t are ranked with respect the frontier and a new population P_{t+1} obtains applying a tournament selection to R_t (see FIG. 2).

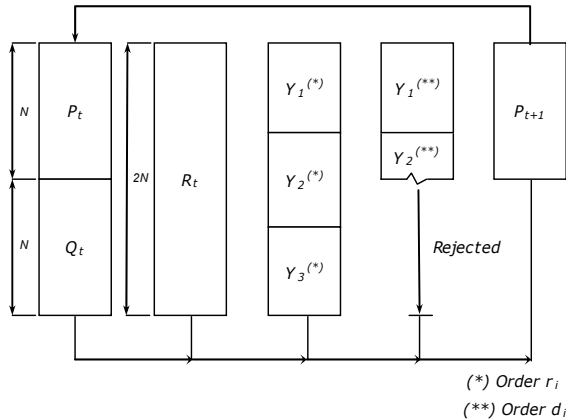


FIG. 2 NON-DOMINATED SORTING GENETIC ALGORITHM II (OUTLINE OF THE MECHANISM OF PROMOTION OF INDIVIDUALS)

After experimenting with several genetic operators we have chosen the Partially Mapped Crossover (PMX) for the crossover and 2-SWAP for mutation. After the individuals have been affected by these operators and before allowing them to become part of a new population we apply an improvement operator. This operator has been designed following the meta-heuristic Simulated Annealing (SA) [Dowland, 1993]. For the change of structure of both chromosomes we select a gen randomly

to change its value. This process, which complements the genetic procedure follows the guidelines of SA.

IV. CONCLUSIONS

We presented a Hybrid Genetic Algorithm (HGA) intended to solve the Flexible Job-Shop Scheduling Problem (FLEX-JSSP). The application of HGA required the calibration of parameters, in order to yield valid values for the problem at hand, which are also a reference for similar problems. We are currently experimenting with other techniques of local search in order to achieve a more aggressive exploration. We are also interested in evaluating the performance of the procedure for other kinds of problems to see whether it saves resources without sacrificing precision in the convergence.

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