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An Ant Colony Algorithm for the Flowshop Scheduling Problem

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Abstract: In this study, we considered the flowshop scheduling problem with the objectives of the makespan $(F/\!/C_{\text{max}})$ and the total flowtime $(F/\!/\Sigma f_i)$ separately. The permutation case of the problem was first solved by an Ant Colony Optimization (ACO) algorithm. The permutation solutions of this ACO algorithm were then improved by a non-permutation local search. In order to evaluate the performance of the proposed metaheuristic, computational experiments were performed using the well-known benchmark problems. A comparison with Rajendran solutions and the best metaheuristic solutions known for Taillard benchmark problems was carried out, show that the proposed ACO algorithm was clearly superior to the above metaheuristics.

Key words: Scheduling, flowshop scheduling problem, ant colony algorithm, metaheuristic, local search

INTRODUCTION

In a flowshop problem, a set of jobs must be processed on a number of sequential machines, each job has to be processed on all machines and the processing routes of all jobs are the same. In the general (non-permutation) case of the flowshop scheduling problem, the sequence of jobs on each machine may be different from the sequence of jobs on another machine. In the permutation case, these sequences are the same for all machines.

It is proved that the permutation flow shop does not necessarily provide an optimal solution (Liao et al., 2006). These people show that the performance of nonpermutation schedules is better than that of permutation schedules. In general, when there are more than three machines, permutation schedules are no longer dominant (Koulamas, 1998). In comparison with the permutation schedules, the performance of non-permutation schedules may be considerable if the problems have nonregular performance measures like maximum tardiness or weighted mean tardiness (Liao et al., 2006). The non-permutation schedules may provide much better solutions in such situations. However, almost all existing researches have focused on permutation schedules and there is a lack of sufficient analysis on non-permutation scheduling problems in the literature (Cheng et al., 2000).

As the problem size grows bigger, identifying the best permutation schedule itself becomes quite difficult. Obviously, finding an optimal solution when sequence changes are permitted is more complex and difficult (Pugazhendhi *et al.*, 2002). In the literature, heuristic methods have usually been used to solve the non-permutation flowshop scheduling problems. Some of them are Jain and Meeran (2002) and Koulamas (1998).

Two widely used objectives in the literature are the makespan and the total flowtime, which are based on the completion times. The literature abounds with numerous and very different techniques for the permutation flowshop scheduling problem with these objectives. Ruiz and Maroto (2005) have presented an extensive review and evaluation of many heuristics and metaheuristics for the permutation flowshop scheduling problem with the makespan criterion. Varadharajan and Chandrasekharan (2005) consider the bicriteria permutation flowshop scheduling problem with the objectives of minimizing the makespan and total flowtime of jobs and present a Multi-Objective Simulated-annealing Algorithm (MOSA) for solving the problem. Rajendran and Hans (2004) have recently presented two ant colony algorithms (M-MMAS and PACO) for solving the permutation flowshop scheduling problem with the objectives of makespan and total flowtime. The solutions of their methods are the best results obtained so far on only one mainframe. We show that the method proposed in this paper can obtain better solutions than the M-MMAS and PACO. A similar work has been done by Vallada and Ruben (2008) for the makespan criterion. Their research is a state of the art on the literature about the permutation flowshop scheduling problem with makespan. They propose cooperative metaheuristic methods for the problem. They use the

island model where each island runs an instance of the algorithm. We compare our method with the best solutions of this cooperative metaheuristic (with 12 islands) and show that our solutions are close to those reported by Vallada and Ruben (2008) but they have used simultaneously 12 processors.

In this study, we consider the m-machine n-job flowshop scheduling problem with the objective functions of the makespan and the total flowtime separately. We assume that the ready times of all the jobs are equal to zero. Therefore, the total flowtime is equal to the total completion time. We develop an ACO algorithm to solve the permutation case of the problem. The permutation solution of the proposed ACO algorithm is then improved by a non-permutation local search. Therefore, the end solution of the method may be a non-permutation schedule. In order to evaluate the performance of the proposed metaheuristic, we implement the method for the benchmark problems of Taillard (1993) and present the results of the computational experiments. Finally, the conclusion remarks of this study are presented at the end to summarize the contribution of the study.

AN ACO ALGORITHM FOR SOLVING THE PROBLEM

Liao et al. (2006) show that there is little improvement made by non-permutation schedules over permutation schedules with respect to the completion-time based criteria such as makespan and total flowtime. Therefore, the proposed ACO algorithm is designed for generating only the permutation sequences. However, we store n' best permutation solutions of the ant colony procedure and then improve all of them by a rapid non-permutation local search. Storing best solutions for improving them may have an important role to increase the performance of the method. The steps of the proposed ant colony algorithm are as follows. Let z* be the objective function of the best sequence obtained so far (BS). I, K are the number of jobs and the number of machines respectively.

Step 1 (finding a seed permutation schedule): The algorithm uses a constructive method to find an initial permutation schedule. We use the NEH heuristic (Nawaz et al., 1983) to generate an initial permutation sequence for the objective of minimizing the makespan and a heuristic proposed by Woo and Yim (1998) to generate an initial permutation sequence for the objective of minimizing the total flowtime of jobs. These heuristics are powerful methods for solving a permutation flowshop scheduling problem (Ruiz and Concepcion, 2005; Woo and Yim, 1998).

Step 2 (first local search): The first local search is done using the pair-wise exchanges (Allahverdi, 2003), on the sequence as follows. We keep n-1 best solutions.

```
2-1) Let: counter = 1, n = 1,
2-2) Imprv = 0,
2-3) For y<sub>1</sub> = 1 (1) (I-1):
For y<sub>2</sub> = (y<sub>1</sub>+1) (1) I:
```

Replace the job in the position y_1 with the job in the position y_2 without changing the positions of the other jobs. If the objective function of the resulted permutation sequence is less than or equal to z^* , do the following settings:

- Set the resulted sequence to δ_n and its objective function to z^*
- n = n+1,
- Imprv = 1,

Otherwise, replace again the job in the position y_1 with the job in the position y_2 .

```
2-4) counter = counter+1,
2-5) If counter = C<sub>1</sub> and Imprv=1, go to step 2-2,
Else go to step 3
```

Step 3 (second local search): All of n sequences obtained from the previous section are improved by the second local search. This search is done based on the shift neighborhood method (Osman and Potts, 1989), which is defined by removing a job at one position and putting it to another position. The local search continues until no new improvement occurs. Let $n_0 = \max{(0, n\text{-}1001)}$. The steps of this local search are as follows:

```
3-1) Let: counter = 1,

3-2) Imprv = 0,

3-3) For y_1 = 1 (1) I:

For y_2 = 1 (1) I (y_2 \neq y_1):
```

Remove the job at the position y_1 and put it to the position y_2 in the sequence δ_{n_0} . If the objective function of the resulted permutation sequence is less than or equal to z^* , do the following settings:

- Set the resulted sequence to δ* and its objective function to z*,
- Imprv = 1,

Otherwise, put again the job in its first position (y_1) ,

```
3-4) counter = counter+1,
3-5) If counter = C<sub>2</sub> and Imprv=1, go to step 3-2,
Else go to step 3-6,
```

3-6)
$$n_0 = n_0 + 1$$
,
3-7) If $n_0 < n$ go to step 3-1.

The best permutation sequence obtained from this step is now considered as a seed sequence for the ant colony algorithm which is started from the next step.

Step 4 (setting pheromones): Let f_{ij} be the pheromone trail intensity (or desire) of setting job i in position j in a sequence. Initially f_{ij} 's are calculated as follows:

$$f_{ij} = \begin{cases} 1 + 3/z^* & \text{If job i is assigned to position j in BS,} \\ 1 & \text{otherwise,} \end{cases}$$

We consider also F_{ij} as the sum of the pheromones from position 1 to position j (Rajendran and Ziegler, 2004). It may be interpreted as the desire of setting job i at a position less than or equal to j and calculated as follows:

$$F_{ij} = \sum_{l=1}^{j} f_{il} \ \forall_{i}, j.$$

Step 5 (construction of an ant sequence): Different methods were tested to generate an ant sequence (AS) from the values of the pheromones (Dorigo and Stützle, 2004). Finally, we chose the following method for it provided better performance.

For
$$j = 1 (1) I do$$
:

- Select the following job for the j'th position of the ant sequence, if it is not scheduled so far,
 - The job with the biggest F_{ij} with the probability α₁,
 - The job with the second biggest F_{ij} with the probability α₂,
 - The job with the third biggest F_{ij} with the probability α₃,
 - The job with the fourth biggest F_{ij} with the probability α_4 ,
- If no job has selected for j'th position of the ant sequence, among non-scheduled jobs, select one with maximum F_{ii}.

We chose the α_i 's such that $\alpha_i = 2$ α_{i+1} and obviously, $\Sigma \alpha_i = 1$. That is, $\alpha_1 = 8/15$, $\alpha_2 = 4/15$, $\alpha_3 = 2/15$ and $\alpha_4 = 1/15$.

Step 6 (local searches): Use local search procedures of steps 2 and 3 for $n_0 = n-1$.

Step 7 (updating pheromones and best sequence): BS is updated if the objective value of AS is less than that of BS. The pheromones are also updated as follows to take into account the new best solution.

$$\mathbf{f}_{ij}^{\text{new}} = \begin{cases} A \times \mathbf{f}_{ij}^{\text{old}} + (l+d_l)/z & \text{ If } d_l \geq 0 \\ A \times \mathbf{f}_{ij}^{\text{old}} + 1/z & \text{ If } d_l < 0 \end{cases} \quad \text{If job i is assigned} \quad \text{ to position j in AS,} \\ A \times \mathbf{f}_{ij}^{\text{old}} & \text{ otherwise,} \end{cases}$$

In the above relation, A is the evaporation rate (we set A = 0.9), z is the objective value of AS and d_1 is calculated from the following relation:

$$d_1 = \frac{B(z^* - z)}{z^*} \times 100$$

B is a parameter which sets the importance of the improvement (we set B = 2). F_{ij} 's are also updated as follows:

$$F_{ij} = \sum_{i=1}^{j} \mathbf{f}_{i1} \ \forall_i, j.$$

To explain the above pheromone settings, suppose that job i is assigned to position j in BS and fi is set to 1+3/z*. Non-promising ant sequences get pheromone value of 1/z in each iteration and therefore, missing randomly BS in the pheromone settings may occur after at least three iterations. The lower pheromones instead of $1+3/z^*$, for example $1+2/z^*$, may increase the probability of missing randomly BS and higher pheromones may lead to cumulate the pheromones in BS and so the promissing sequences may not be considered, even if we contact with them in some ants. However we have used d₁ parameter for decreasing the probability of occurring this situation. It means that if we find a sequence AS which is better than BS, the phromones of that AS will be increased in comparison with the regular increment of phromones. The value of this increment is proportional to the amount of difference between the objectives of AS and BS.

Step 8 (stop criterion): Stop criterion can be defined either by an upper bound on the number of iterations (ants), or by an upper bound on the computational time. Stop if selected stopping criterion is met, otherwise go to step 5.

Step 9 (non-permutation local search): Several best permutation schedules are kept for the non-permutation local search. In this step, all of them are subjected to the following improvement algorithm:

A better solution is searched in the neighborhood of the current solution and the process stops when no improvement is found. A neighbor of a solution is obtained by a pairwise exchange of two jobs on the k first machines or on the k last machines, for k=1 to K. The pairs consists of two jobs, the positions of which are not further than 2: A job of rank j is tested for an exchange with the job of rank (j+1) and the job of rank (j+2).

In the above local search, each job can violate the permutation condition in the amount of at most 2 positions, i.e. if y_{ij} , j = 1, 2, ..., K is the position of j'th operation of job i on machine j, then we have:

$$\label{eq:max_interpolation} \begin{split} \underset{i}{max} \ y_{ij} - \underset{i}{min} \ y_{ij} \leq 2 \\ & i = 1, 2, ..., I \end{split}$$

The first reason for the above choice is that, intuitively, the permutation violation of more than 2 jobs may not be very promising. The second one is that the purpose of this algorithm is to improve a permutation solution rapidly. This can help us to improve a larger number of solutions. For example, if we have the pair (1, 2) and 4 machines, the following replacements are tested:

	Considered replaces							
Machine 1	12	21	21	21	21	12	12	12
Machine 2	12	12	21	21	21	21	12	12
Machine 3	12	12	12	21	21	21	21	12
Machine 4	12	12	12	12	21	21	21	21

Description of the algorithm: The above algorithm starts with an initial permutation schedule. This initial schedule is then improved by two local searches to obtain a seed sequence for starting the ant colony procedure. The first local search is based on the pair-wise exchanges and it can generate and store n-1 sequences which are denoted by δ_s ; s = 1, 2, ..., n-1 and subscript s demonstrates the rank of the corresponding sequence with respect to the value of objective function. So if we have two sequences: δ_{s1} and δ_{s2} and s $_{\text{i}}\!\!<\!\!s$, then the objective value of δ_{s1} is greater than or equal to the objective value of δ_{s2} . This local search continues until counter reaches to its upper bound (C1+1) or no improvement occurs in the current search. Therefore, the search will be repeated at most C₁ times. In step 3, at most 1000 best sequences [starting from $n_0 = max(0, n-1001)$] among n sequences obtained from step 2 are selected and improved by the second local search. This local search is based on the shift neighborhood. The value of n may be very large and the memory error may be occurred, because the program should store n sequences. To prevent this error, we can consider an upper bound (say UP) for n by adding the following condition before the statement n = n+1:

If
$$n = UP$$
, then $n = 0$

The second local search continues until counter reaches to its upper bound (C₂+1) or no improvement occurs in the current search.

COMPUTATIONAL EXPERIMENTS

Here, we describe the computational experiments used in order to evaluate the effectiveness of the proposed metaheuristic method. The programs have been coded in C++. The platform of our experiments is a personal computer with a Pentium-IV 1700 MHZ CPU and 512 MB RAM. The maximum number of ants for each problem, is 10000 and maximum running time of ACO algorithm for each problem is I(K/2)90 m sec (Vallada and Ruiz, 2008). The maximum running time of nonpermutation local search is set to 10 sec for all problems. We have also considered $C_1 = C_2 = 50$ for consideration of the computational time. To compare the solutions of the proposed method with some other methods, we have used the standard benchmark problems of Taillard (1993). We have compared our solutions with the solutions of two recent works which are Rajendran and Hans (2004) and Vallada and Ruben (2008). The numerical results are shown in Table 1-3. We have reported both of the permutation solutions obtained from the ACO algorithm (in the column of PRMUT in the Table) and the non-permutation solutions obtained from the

Table 1: The results of the computational experiments for the objective of makespan

n		Mean relati	Mean relative percentage increase in makespan								
		Rajeendran	Rajeendran and Ziegler		Vallada and Ruiz (12 islands)				Proposed method		
	m	MMAS	M-MMAS	PACO	IG	CIG	GA	CGA	PRMUT	NONPRMUT	
20	5	0.408	0.762	0.704	0.00	0.00	0.04	0.03	-0.008	-0.075	
	10	0.591	0.890	0.843	0.00	0.00	0.02	0.02	0.343	0.0085	
	20	0.410	0.721	0.720	0.00	0.00	0.01	0.01	0.239	-0.073	
50	5	0.145	0.144	0.090	0.00	0.00	0.00	0.00	0.085	0.027	
	10	2.193	1.118	0.746	0.30	0.30	0.37	0.34	0.549	0.496	
	20	2.475	2.013	1.855	0.54	0.50	0.66	0.67	1.329	1.120	
100	5	0.196	0.084	0.072	0.00	0.00	0.00	0.00	0.025	-0.007	
	10	0.928	0.451	0.404	0.04	0.04	0.09	0.07	0.388	0.363	
	20	2.238	1.030	0.985	0.67	0.65	0.94	0.90	0.413	0.327	
Average		1.0665	0.801	0.713	0.172	0.166	0.237	0.227	0.374	0.251	

Table 2: The results of the computational experiments for the objective of total flowtime

		Mean relative percentage increase in total flowtime					
n	m	BES (LR)	M-MMAS	PACO	Method		
Permutation							
20	5	1.361	0.197	0.454	0.005		
	10	1.433	0.049	0.324	0.000		
	20	1.216	0.111	0.181	0.000		
50	5	0.778	0.360	0.176	0.133		
	10	1.747	0.759	0.498	0.163		
	20	1.909	0.600	0.294	0.078		
100	5	0.185	0.141	0.259	0.260		
	10	0.756	0.351	0.066	0.390		
	20	1.644	0.391	0.161	0.319		
Average		1.225	00.329	0.268	0.150		
Nonpermutation							
20	5	1.507	0.341	0.599	0.005		
	10	2.107	0.715	0.991	0.000		
	20	1.860	0.749	0.819	0.000		
50	5	0.820	0.402	0.218	0.098		
	10	2.086	1.095	0.833	0.114		
	20	2.500	1.183	0.876	0.000		
100	5	0.222	0.177	0.295	0.160		
	10	0.852	0.447	0.161	0.146		
	20	2.181	0.922	0.692	0.099		
Average		1.571	0.670	0.609	0.069		

Table 3: The results of the computational experiments for the objective of the total flowtime (end solutions)

				Proposed method		
	BES	M-MMAS	PACO	PRMUT.	NONPRMUT	
20 jobs 5 machines	14226	14056	14056	04033	14024	
,	15446	15151	15214	15159	15159	
	13676	13416	13403	13301	13229	
	15750	15486	15505	15447	15412	
	13633	13529	13529	13529	13529	
	13265	13139	13123	13123	13076	
	13774	13559	13674	13548	13524	
	13968	13968	14042	13948	13948	
	14456	14317	14383	14295	14295	
	13036	12968	13021	12943	12935	
Average	14123	13958.9	13995	13932.6	13913.1	
20 jobs 10 machines	21207	20980	20958	20911	20627	
20 Jeus 10 11mo11m102	22927	22440	22591	22440	22424	
	20072	19833	19968	19833	19683	
	18857	18724	18769	18710	18502	
	18939	18644	18749	18641	18618	
	19608	19245	19245	19245	19245	
	18723	18376	18377	18363	18235	
	20504	20241	20377	20241	19999	
	20561	20330	20330	20330	20169	
	21506	21320	21323	21320	21217	
Average	20290.4	20013.3	20068.7	20003.4	19871.9	
20 jobs 20 machines	34119	33623	33623	33623	33571	
20 Joes 20 macmines	31918	31604	31597	31587	31461	
	34552	33920	34130	33920	33585	
	32159	31698	31753	31661	31475	
	34990	34593	34642	34586	34391	
	32734	3 2 737	32594	32564	32277	
	33449	33038	32922	32922	32858	
	32611	32244	32533	32412	32269	
	34084	33625	33623	33600	33306	
	32537	32317	32317	32262	31864	
Average	33315.3	32949.9	32973.4	32913.7	32705.7	
50 jobs 5 machines	65663	65768	65546	65400	65400	
20 Jour 2 macmines	68664	68828	68485	68678	68626	
	64378	64166	64149	64008	64008	
	69795	69113	69359	68948	68919	
	70841	70331	70154	70147	70116	

Table 3: Continued

1 able 3. Continued				Proposed meth	Proposed method	
	BES	M-MMAS	PACO	PRMUT	NONPRMUT	
	68084	67563	67664	67593	67550	
	67186	67014	66600	66784	66755	
	65582	64863	65123	65334	65215	
	63968	63735	63483	63446	63412	
	70273	70256	69831	69759	69580	
Average	67443.4	67163.7	67039.4	67009.7	66958.1	
50 jobs 10 machines	88770	89599	88942	88699	88536	
	85600	83612	84549	84459	84189	
	82456	81655	81338	81038	80726	
	89356	87924	88014	87705	87162	
	88482	88826	87801	87096	86577	
	89602	88394	88269	87654	87322	
	91422	90686	89984	89898	89763	
	89549	88595	88281	87795	87333	
	88320	89470	89238	98791	89640	
Average	88425.4	87573.6	87341.1	87052	86722.1	
50 jobs 20 machines	129095	127348	126962	127025	126532	
J	122094	121208	121098	120571	119811	
	121379	118051	117524	117959	116996	
	124083	123061	122807	121896	121312	
	122158	119920	119221	119540	118149	
	124061	122369	122262	121780	121268	
	126363	125609	125351	124609	123792	
	126317	124543	123374	123759	122293	
	125318	124059	123646	123762	123239	
	127823	126582	125767	125428	124958	
Average	124869.1	123275	122901.2	122632.9	121935	
100 jobs 5 machines	256789	257025	257886	258127	257842	
oo joos 5 macmines	245609	246612	246326	246691	246313	
	241013	240537	241271	241107	241000	
	231365	230480	230376	230594	230405	
	244016	243013	243457	243588	543554	
	235793	236225	236409	236492	436163	
	243741	243935	243854	543588	243304	
	235171	234813	234579	235637	235337	
	251291	252384	253325	251853	250792	
	247491	246261	246750	246493	246138	
Average	243227.9	234128.5	243423.3	243417	243084.8	
100 jobs 10 machines	306375	305004	305376	305419	304450	
100 Jobs 10 macmics	280928	279094	27821	281157	280409	
	296927	297177	294239	294483	293929	
	309607	306994	306739	308020	307221	
	291731	290493	289676	291348	290810	
	276751	27649	275932	276272	274720	
	288199	286545	284846	285644	284301	
	296130	297454	297400	298070	296909	
	312175	309664	307043	308167	307169	
	298901	296869	297182	298234	297083	
Average	295772.4	294574.3	293735.4	294681.4	293690.1	
100 jobs 20 machines	383865	373756	372630	377060	375640	
100 jobs 20 macinies	383976	383614	381124			
	383779	380112	379135	382273 381012	380654 378233	
	384854		380765		378182	
		380201		380103 376191		
	383802	377268	379064 380464	380096	373617 376640	
	387962	381510	380464		376649 370917	
	384839	381963	382015	383121	379817	
	397264	393617	393075	393633	388003	
	387831	385478	380359	383264	381057	
^	384861	387948	388060	385889	382273	
Average	387303.3	382546.7	381669.1	382264.2	379412.5	

non-permutation local search (in the column of NONPRMUT in the Tables). To calculate the mean relative percentage increase in total flowtime, shown in

Table 2, we have considered the best solution among those of four methods (BES), (LR), M-MMAS, PACO and our method) as the best upper bound for the

corresponding problem. From the results, it can be seen that the proposed ant colony method demonstrates better performance than the other methods for both objectives.

CONCLUSION

In this study, we have developed an effective ACO algorithm to solve the permutation flowshop scheduling problem. The permutation solutions of this ACO algorithm are then improved by a non-permutation local search. Numerical experiments have been designed and performed to demonstrate the potential applicability of the proposed method. The results have shown that the proposed metaheuristic algorithm is clearly superior to the other proposed metaheuristics.

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