

Design of a Hybrid Genetic Algorithm for Parallel Machines Scheduling to Minimize Job Tardiness and Machine Deteriorating Costs with Deteriorating Jobs in a Batched Delivery System

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Abstract

This paper studies the parallel machine scheduling problem subject to machine and job deterioration in a batched delivery system. By the machine deterioration effect, we mean that each machine deteriorates over time, at a different rate. Moreover, job processing times are increasing functions of their starting times and follow a simple linear deterioration. The objective functions are minimizing total tardiness, delivery, holding and machine deteriorating costs. The problem of total tardiness on identical parallel machines is NP-hard, thus the under investigation problem, which is more complicated, is NP-hard too. In this study, a mixed-integer programming (MILP) model is presented and an efficient hybrid genetic algorithm (HGA) is proposed to solve the concerned problem. A new crossover and mutation operator and a heuristic algorithm have also been proposed depending on the type of problem. In order to evaluate the performance of the proposed model and solution procedure, a set of small to large test problems are generated and results are discussed. The related results show the effectiveness of the proposed model and GA for test problems.

Keywords: Parallel machine scheduling, Machine deterioration, Job deterioration, Batched delivery system, Genetic algorithm.

1. Introduction

Parallel machine scheduling as a very traditional branch of research studies have addresses from various aspects with an endeavor to minimize makespan, tardiness, completion time, waiting time, idle time and so on (Bandyopadhyay & Bhattacharya, 2013). In a real world application, the performance of each machine deteriorates over time at a random rate caused by processing the jobs. Moreover, while waiting for processing, jobs also may deteriorate. Deterioration of a job means that a job processing time is defined by a function of its starting times and positions in the sequence (Mazdeh et al., 2010).

On the other hand, manufacturers aims at schedule their production activities such that be able to meet the due dates of their customers' orders. The other issue accounted for in scheduling is the delivery costs of the jobs or Designing a batch of products, to be delivered. Batched delivery system means that the jobs of each customer could be delivered to it in one or many batches. Such a delivery system may leads to different completion and delivery/rendition times for

jobs. Although batched delivery system seems a reasonable approach in reducing transportation costs, but it may leads to increasing the number of tardy jobs or the total tardiness (Hamidinia et al., 2012; Karimi & Davoudpour, 2015).

The problem of scheduling with the batch delivery system first introduced by Cheng and Kahlbacher (Cheng & Kahlbacher, 1993). They studied the single-machine batch delivery scheduling problem with the objective of minimizing the sum of earliness penalties, as well as delivery costs. Also, the authors proved that the problem is NP-hard in the ordinary sense but polynomially solvable for equal weights. In another study, Wang and Cheng (Wang & Cheng, 2000) considered a problem of parallel-machine batch delivery scheduling with the objective of minimizing the sum of flow times and delivery cost and showed that the problem is strongly NP-hard. They developed dynamic programming algorithm for the problem and presented two polynomial time algorithms in the cases that the job assignment is predetermined or the job processing times are all identical.

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Hall and Potts (Hall & Potts, 2003) provided dynamic programming solutions for a range of scheduling problems with batched delivery systems in a supply chain with the aim of minimizing overall scheduling and delivery costs, cost, using several classical scheduling objectives. Mazdeh et al. (Mazdeh et al., 2007) proposed branch and bound algorithms for weighted sum of flow times in a batched delivery system, when all jobs are available at the time zero and in the presence of ready time. Mazdeh et al. (Mazdeh et al., 2011) developed a branch-and-bound algorithm for single machine batch delivery scheduling problem with the objective of minimising the sum of weighted flow times and delivery costs and with the assumption that the delivery cost is a linear increasing function of the number of deliveries.

Hamidinia et al. (Hamidinia et al., 2012) suggested a genetic algorithm for single-machine batch delivery scheduling problem with the objectives of minimizing the sum of earliness, tardiness, holding, and delivery costs. Moreover, the authors presented a mathematical model for the problem. Yin et al. (Yin et al., 2014) addressed the problem of scheduling n nonresumable and simultaneously available jobs on a single machine in which the jobs were delivered in batches to the customers. They assumed a fixed unavailability interval for the machine during which the production is not allowed and along with a scheduling decision, a batching decision has to be determined so as to minimize the sum of total flow time and batch delivery cost, where the cost per batch delivery is fixed and independent of the number of jobs in the batch. They showed that the problem is NP-hard based on a reduction from the Equal-Size Partition Problem and presented a pseudo-polynomial time dynamic programming algorithm. Moreover, they developed a fully polynomial time approximation scheme (FPTAS) and a bicriteria fully polynomial time approximation scheme.

Recently, Ahmadizar and Farhadi (Ahmadizar & Farhadi, 2015) addressed a single-machine batch delivery system scheduling problem to minimize the sum of earliness, tardiness, holding, and delivery costs in which jobs were released in different points in time. They presented a mathematical model and a set of dominance properties. In order to solve this NP-hard problem, they proposed a hybrid algorithm by integrating the dominance properties with an imperialist competitive algorithm. Karimi and Davoudpour (Karimi & Davoudpour, 2015) addressed the scheduling of supply chain with interrelated factories containing suppliers and manufacturers in which jobs transportation among factories and also delivery to the customer can be performed by batch of jobs. In order to

solve the problem, branch and bound algorithm was adopted and a lower bound and a standalone heuristic which was used as an upper bound were also introduced.

On the other hand, Mazdeh et al. (Mazdeh et al., 2010) studied the problem of parallel machines scheduling to minimize job tardiness and machine deteriorating cost with deteriorating jobs and proposed LP-metric method to show the importance of their proposed multi-objective problem. It should be noted that the main difference of our study by Mazdeh et al. (Mazdeh et al., 2010) is that we consider batch delivery system for the same parallel machine scheduling while in Mazdeh et al. (Mazdeh et al., 2010) there is no assumption of batch scheduling problem. Moreover, in addition to machine deteriorating cost with deteriorating jobs which are addressed in both study, because of the inherent properties of the batch delivery scheduling problem, objectives of delay, holding (inventory) and delivery costs are added.

Although batch scheduling problem have already been studied in a number of studies, but most of them have addressed the concerned problem in the case of single machine. Moreover, there is no existing study which proposes a mathematical model for the parallel machines batch delivery scheduling problem in the deteriorating environment.

This paper aims at solving the problem of parallel machine scheduling to minimize total tardiness costs of deteriorating jobs plus their delivery costs and machine deteriorating costs in a batched delivery system when all jobs are available at time zero and preemption is not allowed. In order to solve the presented model, a hybrid genetic algorithm is proposed. To the best of our knowledge, there is not any paper that addresses the assumptions which are considered in this paper simultaneously.

The rest of this paper is organized as follows: section 2 provides a review on the related literature. Section 2 defines the problem and presents the mathematical model for the problem. The proposed hybrid GA is developed in section 3, and finally Section 4 describes the computational results.

2. Problem description

2.1. Problem definition

There are N independent jobs has to be processed on m parallel machines. Machines can process all jobs in different speeds. The jobs belong to F customers while each of them has $o(j)$ orders/jobs ($1 \leq j \leq F$). The processing time of the i th job of the j th customer on machine m is given as a linear increasing function dependent on its starting time and a

constant part ($p_{ijm}=a_{ijm}+b_{ij} s_{ijm}$). a_{ijm} and b_{ij} denotes the constant part and deterioration rate of job i of customer j . constant part of processing time of a job (a_{ijm}) varies with different machine. The deterioration rate of the processing time (b_{ij}) is independent of machine. All jobs are available at time zero, no preemption is allowed, and the setup time on each machine is included in the constant part of the processing time.

Deterioration effect occurred on both job and machine; but deterioration of jobs and machines are independent. Deterioration cost of a machine (M_{ijm}) is dependent on both the machine and the job processed on that machine.

Each job has a completion time c_{ij} , a delivery time or due date d_{ij} , a rendition time R_{ij} , a delay cost α_{ij} , a holding cost h_{ij} , a delivery cost D_j and machine deteriorating cost. A job is tardy when its rendition time is after its delivery time or due date ($R_{ij}>d_{ij}$) and it is called on time when its delivery time equal to its due date ($R_{ij}=d_{ij}$).

Each job can be submitted to the customer when it is processed or its delivery can be postponed to be conducted in a batch of other jobs from the same customer. The rendition time of a job is then equal to the rendition time of the batch containing that job, because the delivery of all jobs of a batch is postponed to the completion time of the last job of the current batch. (Hamidinia et al., 2012)

We consider three types of cost in the studied problem. First type is related to penalty costs. Second type is related to batched delivery system and third type is the cost related to the deteriorating of the machines. Penalty costs include holding costs and delay costs. Holding occurred when the job is completed before it is rendered ($C_{ij}<R_{ij}$). Delay cost occurred when the job is delivered after its due date ($d_{ij}<R_{ij}$). So tardiness of a job (T_{ij}) and holding of a job (H_{ij}) are as follows:

$$T_{ij} : \max \{0, (R_{ij} - d_{ij})\}$$

$$H_j : R_{ij} - C_{ij}$$

According to the common form for classifying scheduling problems suggested by graham et al. (Graham, Smiley, Russell, & Nairn, 1977), the problem can be summarized as

$1 \parallel \sum C_{ij} + \sum H_j + \sum D_j + \sum M_{ijm} \mid s_{ijm} < \infty$

1. Minimizing the sum of delay, holding (inventory), delivery and machine deteriorating costs
2. Finding the optimal number of batches and assignments of jobs to the batches
3. Finding the optimal assignment of jobs to machines

2.2. Notations

Decision variables and parameters used in the mathematical model are as follows:

Indices

j	customers, $j = 1, \dots, F$
i	jobs/orders which are related to a customer, $i = 1, \dots, o_j$
m	machines, $m = 1, \dots, M$
k	batches, $k = 1, \dots, K$
e	positions on machines, $e = 1, \dots, E$

3 D U D P H W H U V

1 W R W D O Q X P E H U R I P D F K L Q H V D Y D
 E D W (A K H V L O)

0 W R W D O Q X P E H U R I P D F K L Q H V D Y D
 R W K H Q X P E H U R I R U G H U V Z K L F K U H
 . L M G H O D \ F R V W R I W K H L W K M R E R I W
 K L M K R O G L Q J F R V W R I M R E L R I W K H F
 D L M P F R Q V W D Q W S D U W R I W K H S U R F H V
 F X V W R I P R I Q P D F K L Q H P
 E L M G H W H U L R U D W L Q J U D W H M R I M R E L R
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 E L J O S O D U J H S R V L W L Y H Q X P E H U

Decision variables

X_{ijk}	a binary variable which indicates the assignment of the i th job of the customer j to k th batch
X_{ijek}	a binary variable which indicates the assignment of the i th job of the customer j to e th part of k th batch
Y_{jk}	a binary variable indicating whether there is a job relates to the customer j which is assigned to k th batch
τ_{ij}	a binary variable which indicates whether or not the i th job of the customer j is tardy
Z_{ijm}	a binary variable which indicates the assignment of the i th job of the customer j to machine m
W_{ijlm}	a binary variable which indicates the assignment of the i th job of the customer j on situation l at machine m

P_{ijm} processing time of the i th job of the customer j on machine m
 S_{ijm} starting time of job i of the customer j on machine m
 $C_{L M}$ completion time of job i of the customer j
 C_{k} completion time of k th batch
 $C_{L M}$

T_{ij} tardiness of job i of the customer j

2.3. Mathematical model

According to the above mentioned description of the problem, a mathematical model is proposed based on (Hamidinia et al., 2012; Mazdeh, Rostami, & Namaki, 2013; Mazdeh et al., 2010) which is as follows:

$$C_{ij} = P_{ijm} + S_{ijm} + T_{ij} \quad \forall i, j, m$$

$$C_{k} = C_{L M} + C_{k} \quad \forall k$$

$$C_{L M} = C_{k} + C_{L M} \quad \forall k, m$$

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(13)

$$C_{ij} \geq E_{ij} + C_{ij} \cdot \delta_{ij} \quad (14)$$

$$C_{ij} \geq C_{ij} \cdot \delta_{ij} \quad (15)$$

$$C_{ij} \geq C_{ij} \cdot \delta_{ij} \quad (16)$$

$$C_{ij} \geq C_{ij} \cdot \delta_{ij} \quad (17)$$

$$C_{ij} \geq C_{ij} \cdot \delta_{ij} \quad (18)$$

$$C_{ij} \geq C_{ij} \cdot \delta_{ij} \quad (19)$$

Objective (1) minimizes the sum of delay, holding, delivery and machine deterioration costs. Equation (2) computes the amount of tardiness for the *i*th job of the customer *j*. Constraint (3), (4) guarantees that a job is tardy if its rendition time is greater than its due date. Constraint (5) computes the completion time of each batch that is equal to the sum of the processing times of all jobs located in the current batch plus the completion time of the preceding batch. Constraint (6) calculates the completion time of the *i*th job related to customer *j* which is equal to the completion time of the preceding batch plus processing times of this job and all the jobs assigned to the current batch before it. Constraint (7) ensures that each job is only assigned to one batch. Constraint (8) guarantees that jobs belonging to different customers not to be batched together i.e., each batch is assigned to the jobs of a one customer. Constraints (9), (10) ensures that a batch is not be assembled until at least one job is assigned to it. Constraint (11) calculates the rendition time of each job which is equal to the completion time of the pertinent batch. Constraint (12) defines the existing parts of each machine.

Constraint (13) states the relation between the processing time, start time and fixed part of processing time of the *i*th job of the *j*th customer which is assigned to machine *m*. Constraint (14) ensures that the starting time of a job on a machine is at least equal to the completion time of the preceding job. Constraint (15) guarantees that each job is assigned to only one of the existing positions on the machines while restriction (16) ensures that on each existing position, at most one job can be assigned. Constraint (17) guarantees that jobs are not allowed to assign to one empty position of a machine until the previous position have been assigned. Constraint (18) defines the existing positions of each machine. Constraint (19) introduces binary and continuous variables.

As it can be seen from above-mentioned model, the resulting model is not linear because variables are multiplied in constraints (2), (5), (6) and (11) and for example $C_{ij} \cdot \delta_{ij}$ and $C_{ij} \cdot \delta_{ij}$ in constraint (2) and $C_{ij} \cdot \delta_{ij}$ in constraint (5). In order to convert the non-linear constraint (5) to the linear form, we use an auxiliary variables and constrains which are defined as follows:

$$C_{ij} \geq E_{ij} + C_{ij} \cdot \delta_{ij} \quad (5-1)$$

$$C_{ij} \geq C_{ij} \cdot \delta_{ij} \quad (5-2)$$

$$C_{ij} \geq C_{ij} \cdot \delta_{ij} \quad (5-3)$$

$$C_{ij} \geq C_{ij} \cdot \delta_{ij} \quad (5-4)$$

$$\sum_{i \in P} Q_i > E \text{ Cál } \sum_{i \in P} \hat{E} \text{ F} \text{ Cál} \tag{5-5}$$

$$\sum_{i \in P} Q_i \sum_{i \in P} \hat{E} \text{ F} \text{ Cál} \tag{5-6}$$

$$\sum_{i \in P} R_i \sum_{i \in P} F_i > E \text{ Cál } \sum_{i \in P} F_i : \sum_{i \in P} \hat{E} \text{ F} \text{ Cál} \tag{5-6}$$

Similarly, for constraints (2), (6) and (11) following constraints are defined:

$$\sum_{i \in L} i_{E \text{ F}} \hat{E} \text{ F} \tag{2-1}$$

$$\sum_{i \in Q} Q_i > E \text{ Cál } \hat{E} \text{ F} \tag{2-2}$$

$$\sum_{i \in Q} Q_i \hat{E} \text{ F} \tag{2-3}$$

$$\sum_{i \in R} R_i \hat{E} \text{ F} > E \text{ Cál } \sum_{i \in R} F_i \hat{E} \text{ F} \tag{2-4}$$

$$\sum_{i \in L} i_{E \text{ F}} \hat{E} \text{ F} \tag{2-5}$$

$$\sum_{i \in Q} Q_i \hat{E} \text{ F} \tag{2-6}$$

$$\sum_{i \in R} R_i \hat{E} \text{ F} > E \text{ Cál } \sum_{i \in R} F_i \hat{E} \text{ F} \tag{2-7}$$

$$\sum_{i \in L} i_{E \text{ F}} \hat{E} \text{ F} \tag{2-5}$$

$$\sum_{i \in L} L_i \hat{E} \text{ F} \tag{2-5}$$

$$\sum_{i \in L} L_i \hat{E} \text{ F} \tag{2-5}$$

$$\sum_{i \in L} L_i \hat{E} \text{ F} \tag{2-5}$$

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$$\sum_{i \in L} L_i \hat{E} \text{ F} \tag{2-5}$$

$$\sum_{i \in L} L_i \hat{E} \text{ F} \tag{2-5}$$

$$4 \sum_{i \in I} \sum_{j \in J} x_{ij} \leq \sum_{k \in K} C_k \quad (11-1)$$

$$\sum_{i \in I} x_{ij} \leq C_j \quad \forall j \in J \quad (11-2)$$

$$\sum_{j \in J} x_{ij} \leq C_i \quad \forall i \in I \quad (11-3)$$

$$\sum_{i \in I} \sum_{j \in J} x_{ij} \leq \sum_{k \in K} C_k \quad (11-4)$$

By replacing constraints (2), (5), (6) and (11) with above mentioned numbered constraints the non-linear model is converted to an equivalent MILP model.

3. GA Implementation

Evolutionary algorithms (EAS) are the most studied population based metaheuristics which are applied to many real and complex problems (Talbi, 2009). GA is a very popular class of EAs which is inspired by the natural evolution of the living organisms. The basic concepts of GA have been described by the investigation studied by John Holland in the 1970s and Davis was first that proposes GA for solving scheduling problems (Davis & Coombs, 1987).

A generic GA starts by creating an initial population of chromosomes (solutions) and iteratively replaces the current population by a new population. Each chromosome/individual encodes a solution of the problem, and its fitness value represents a measure of the quality of solution for the problem. In order to search potential better solutions, during each iteration (generation) genetic operators that are, crossover, mutation and natural selection are applied. Crossover operator creates new individuals by combining parts of two individuals while mutation operator creates new individuals, by a small change in a single individual. The following subsections present the components of the hybrid GA applied to solve the proposed model.

3.1. Solution representation

The initial and most important step in the GA is to consider a chromosome structure or solution representation. In the studied problem, a chromosome structure should contains the following characteristics:

- Assembled batches
- Jobs belonging to each assembled batch
- Order of completing jobs
- Assignment of jobs to machines and order of processing

- comply the restriction of placing of only one customer's jobs in each assembled batch

Hence, we use the chromosome structure presented in Figure 1. For the solution of the presented model. On the other hand, to access the job indices easier, we assign a unique index for each job (j indicates job number) by using orders vector. O(c) is the orders vector of customers; for example o(c) = [2 3 1 2 1] means that there are 5 customers in which first customer has 2 orders/ jobs (o₁=2) and jobs 1 and 2 belong to first customer. Second customer has 3 jobs (o₂=3) and jobs 3, 4 and 5 belong to second customer and so on.

As it depicted in Figure 1 the chromosome structure in this study consists of 4 rows and N (total number of jobs) columns in which the index of columns represent the job number. First row of the chromosome represents the assignment of the jobs to machines. Each gene value in the first row refers to a machine number. Similar to the chromosome structure proposed by hamidinia et al. (Hamidinia et al., 2012), Second row is a permutation of N initial batches that represents the initial batches assigned to the jobs in which the value of each gene is the number of the batch number. The total number of the initial batches is equal to total number of jobs. This row insures that particular batches can be assigned to the jobs of each customer.

First we suppose N empty batches with numbers 1 to N where each number showing the position of corresponding batch in the sequence of batches. i.e., a batch with lower number is processed and delivered before the subsequent batches. Third row of the chromosome represents the real batch number for each customer so repeated numbers are allowed (Some jobs of one customer may place in the same batch). The subsection of each customer in the third row is a subset of related subsection in the second row. The length of each subsection equals the number of orders of the corresponding customer (for example if the first customer has 2 jobs/orders, then the values of the first 2 cells of the

initial batches row could be used for packing this customer's orders).

The 4th row of the proposed chromosome is order priority row that consists of subsections of customer. Each subsection of the order priority row is a permutation of numbers between 1 and the length of the subsection, i.e.,

values in each subsection represents the processing order of related customer's jobs. If some orders of a one customer place in different batches, the processing order of them are based on priority of their batches (according to batch numbers), but if these orders fall in the same batch the priority order specify the order of processing.

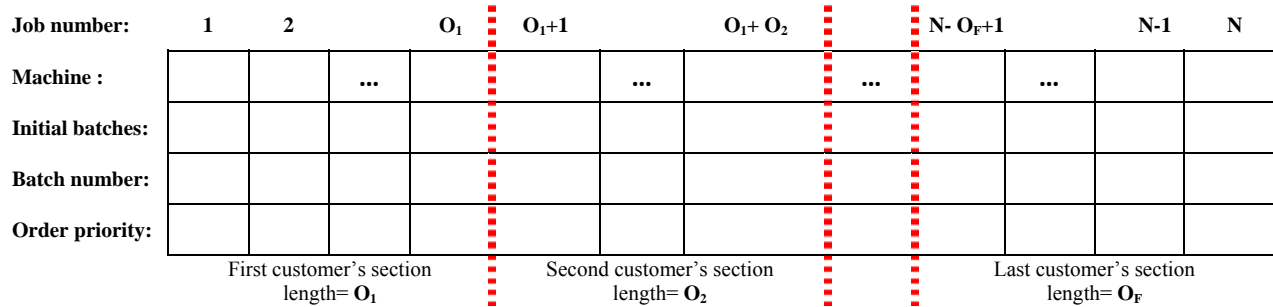


Fig 1. Proposed chromosome structure

3.2. Chromosome decoding and fitness function

In order to interpret a chromosome and construct a solution, first we refer to the batch number row to determining the used batch sets and remove the empty batches from the batch sets. Table 1 depicts the input data of a sample problem with 9 jobs, 5 customers and 3 machines; given orders vector is $o(c)=[2\ 3\ 1\ 2\ 1]$ and j is an index for job number. Figure 2 represents one feasible solution for the problem. By sorting the number of occupied batches we gain the sequence of processing the batches. For example in Figure 2 batch sets with respect to the order of processing them are 1-2-3-5-6-8 (regarding batch number row) which means batches 2, 4, 7 and 9 are empty. For determining corresponding jobs of each batch and each machine we select the first batch and find the jobs related to it. In example of Fig 2. first batch is batch number 1. So we should refer to the *batch number row* to find the cells with value 1. The found cells are 4 and 5 which means job 4 and 5 (related to customer 2) are in the batch number 1. Also for finding the processing order of jobs 4 and 5, we should refer to corresponding cells (cells 4 and 5) in *order priority row* that shows that the order priorities of jobs 4 and 5 are 1 and 2 which means first job to be processed is job 4 & second job is job 5.

Now for finding machine number for job 4 (first job) we consider the cell number 4 in the *machine row* (first row) that shows machine number 3 for job 4. Since job 4 is first job, all the machines are free. After assigning job 4 to machine 3, cell 5 in the machine row indicates that job 5 is assigned to machine 3 too. So the starting time of job 5 is

equal to completion time of job 4 on machine 3. Now that the first batch and its related jobs are scheduled, next batch (batch 2) and its related jobs (jobs 1 & 2) are scheduled and so on. Finally the resulted assignment of jobs to machines and batches determines the order of completing the jobs and delivering the batches. Consequently, by calculating the rendition & completion time and comparing with delivery time; delay, holding, delivery, and machine deteriorating costs can be obtained and sum of these costs is the fitness of the current individual. Table 2 provides the fitness of the chromosome of Fig 2. .

3.3. Crossover

Based on the structure of the proposed chromosome, a two stage approach for crossover has been designed in this paper. In the first stage, crossover is used for *machine row* and *order priority row* and at the second stage a crossover is performed for *initial batches row* & *batch number row*. For the first stage, an extension of uniform crossover is proposed which refer to it as "*customer transfer*" method. The customer transfer method is shown in Fig 3. The algorithmic structure of this method is as follows:

- Randomly generate a binary vector r with length of orders vector (i.e., the length is equal to number of customers). First element of r is related to customer one, second element is related to customer number 2 and so on.

x , I WKHY D ~~OR~~ ~~WKH~~ IWKHOHPHQW
 (sQBQY}RIULV WKHQIFURVSW W
 DQG U^GURZJHOMW ~~W~~ BSKLUREQGRUGHU
 SULRW ~~W~~ ~~W~~ VHFVWLR ~~Q~~ ~~K~~ VWR ~~R~~ ~~R~~ ~~R~~

SDUHQW WR WKH FRUUHVSRQGLQJ ORFDWLRQ RI the initial batches row and then in
 DQG FRS\ WKH JHQHURZ RI VHFVLRQ to preserve the feasibility of the solution, generate
 IURP SDUHQW FRUUHVSRQGLQJ ORFDWLRQ the new batch number row of the resulted offsprings
 RIIVSULQJ from initial batches row. Order crossover can be
 summarized as follows:
 x , IWKHY D WKHIWKHOHPHQW First, two crossover points are randomly selected. From
 (sQBQ) RI U LV WKHQ IURVW WKH first crossover point will copy its offspring
 DQG UG URZ RI VHFVLRQ I IWK FXVWRU RI parent 1 to the offspring, at the same
 WR WKH FRUUHVSRQGLQJ ORFDWLRQ position, the part between the two
 FRS\ WKH JHQHURZ RI VHFVLRQ I IWK FXVWRU For parent 2, one will start at the second crossover
 SDUHQW WR WKH FRUUHVSRQGLQJ ORFDWLRQ point and pick the elements that are not already selected
 from parent 1 to fill them in the offspring from the
 second crossover point. (Talbi, 2009) Crossover stages
 for the solutions of Fig 3 is illustrated in Fig 4.

For applying crossover in the second stage for initial batches row and batch number row, first we use order

Table 1
Input data for example problem

Number of jobs=9; number of customers=5; number of machines=3; o(c)=[2 3 1 2 1]									
Customer NO.	1	2	3	4	5	6	7	8	9
Job No.	1	2	3	4	5	6	7	8	9
Delivery time	12	20	26	20	40	10	30	50	39
Delay cost	10	15	15	12	13	10	15	12	10
Holding cost		5		5		5		5	5
Delivery cost		60		80		40		80	60
fixed part of the processing time(a _m) and growth rate of the processing time(b _j)									
Job No.	1	2	3	4	5	6	7	8	9
Machine 1	10	12	15	5	9	10	12	12	14
Machine 2	12	8	16	7	6	9	10	9	12
Machine 3	14	14	12	9	7	9	13	8	16
b _j	0.35	0.25	0.50	0.25	0.60	0.70	0.20	0.30	0.40
machines deteriorating cost (M _m)									
Job No.	1	2	3	4	5	6	7	8	9
Machine 1	2	3	4	4	5	6.5	8	9	9.5
Machine 2	1.5	3	4.5	4.5	5.5	7	8	8.5	9
Machine 3	2.5	3.5	5	5	6	7.5	8.5	9.5	10

3.4. Mutation

In this study we utilize a mutation operator for machine row, initial batches and priority order rows.

3.4.1. Mutation operator for machine row

In this paper, two types of mutation is performed on the machine row. First type is called "job transfer" in which we select a random gene in the machine row (machine number) and change the value of it to another possible value from the machine set. This operation is equivalent to transfer one job from current machine to another machine. Second type is

swap mutation operator in which the value of two randomly selected genes is replaced. Number of mutation operation is performed (i.e., n(m)) in the machine row is dynamic and dependent on length of a chromosome and number of machines:

$$J: l; L \left\{ \begin{matrix} PKP=H JQI>AN KB FK>O \\ JQI>ANI=?DEJAO^S \end{matrix} \right. : s \{;$$

If n(m) is even, number of "job transfer" and swap processes are equal. And if n(m) is odd, number of swap operations is one unit more than "job transfer" operations. For the sample problem of Fig 2. n(m) is equal to 2 which means that one "job transfer" and one swap move is performed.

Representation of one possible solution:

Job number:	1	2	3	4	5	6	7	8	9
Machine:	2	1	1	3	3	2	3	2	1
Initial batches:	7	2	3	1	9	5	8	4	6
Batch number:	2	2	3	1	1	5	8	8	6
Order priority:	2	1	3	1	2	1	2	1	1

Customer. 1
C. 2
C. 3
C. 4
C. 5

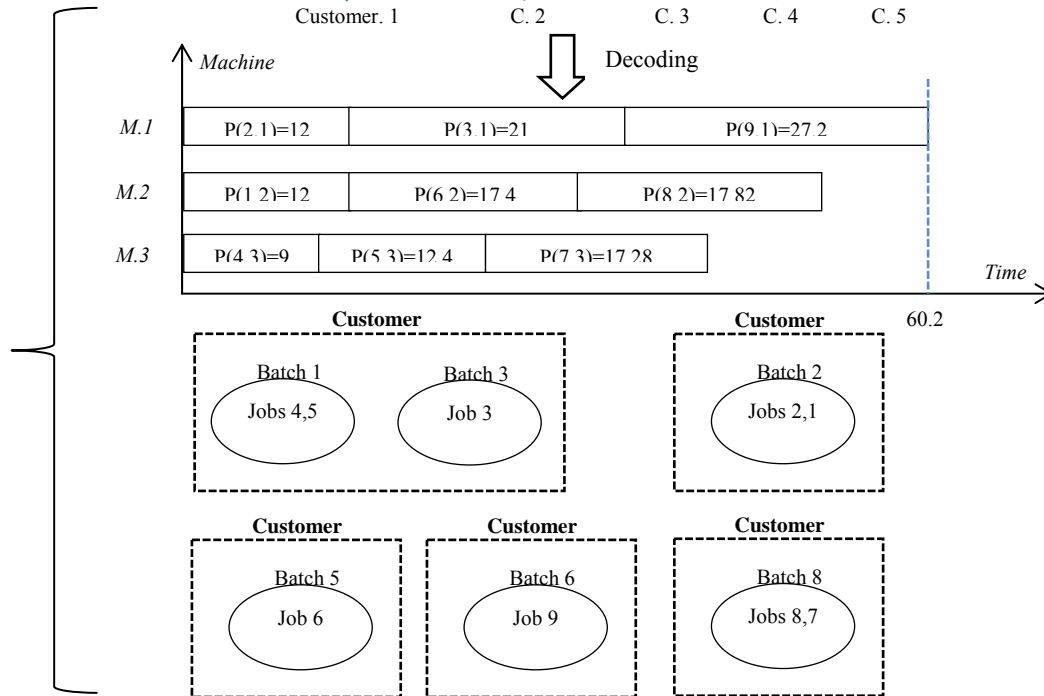


Fig 2. Chromosome decoding for sample problem

Table 2
Fitness function calculations for example problem

Customer NO.	1		2		3	4		5	
Job No.	1	2	3	4	5	6	7	8	9
Batch No.	2	2	3	1	1	5	8	8	6
Order priority	2	1	3	1	2	1	2	1	1
Process order	1	1	2	1	2	2	3	3	3
Machine	2	1	1	3	3	2	3	2	1
Process time	12.00	12.00	21.00	9.00	12.40	17.40	17.28	17.82	27.20
Completion time	12.00	12.00	33.00	9.00	21.40	29.40	38.68	47.22	60.20
Rendition time	12.00	12.00	33.00	21.40	21.40	29.40	47.22	47.22	60.20
Due date	12	20	26	20	40	10	30	50	39
Delay cost	0	0	105.00	16.80	0	194.00	258.30	0	212.00
Holding cost	0	0	0	62.00	0	0	42.70	0	0
Delivery cost	30	30	80	40	40	40	40	40	60
Machine deteriorating cost	1.50	3.00	4.00	5.00	6.00	7.00	8.50	8.50	9.50
Total cost	31.5+33+189+123.8+46+241+349.5+48.50+281.5=1343.8								

3.4.2. Mutation operator for initial batches row

Swap mutation operator is utilized for initial batches row while the number of swap moves is equal to $n(m)$. To preserve feasibility of the batch number row, after applying swap mutation on the initial batches row, new batch number row is generated from the resulted initial batches row.

3.4.3. Mutation operator for order priority

Based on proposed chromosome structure, swap operator for order priority row is done dependently in each customer section of the row which means for each customer. If the number of customers' orders is O_f then number of

customer's orders. Fig. 5 illustrates mutation operator for sample chromosome.

3.5. Proposed heuristic algorithm

A heuristic algorithm is proposed to improve the solutions after mutation. Every mutated solution is improved through the proposed heuristic algorithm. The steps of the proposed algorithm are as follows:

Step1: change the priority order of each customer's orders based on EDD rule.

Step 2: assign two consecutive jobs of each batch to different machines.

Step 3: assign the jobs uniformly to the machines.

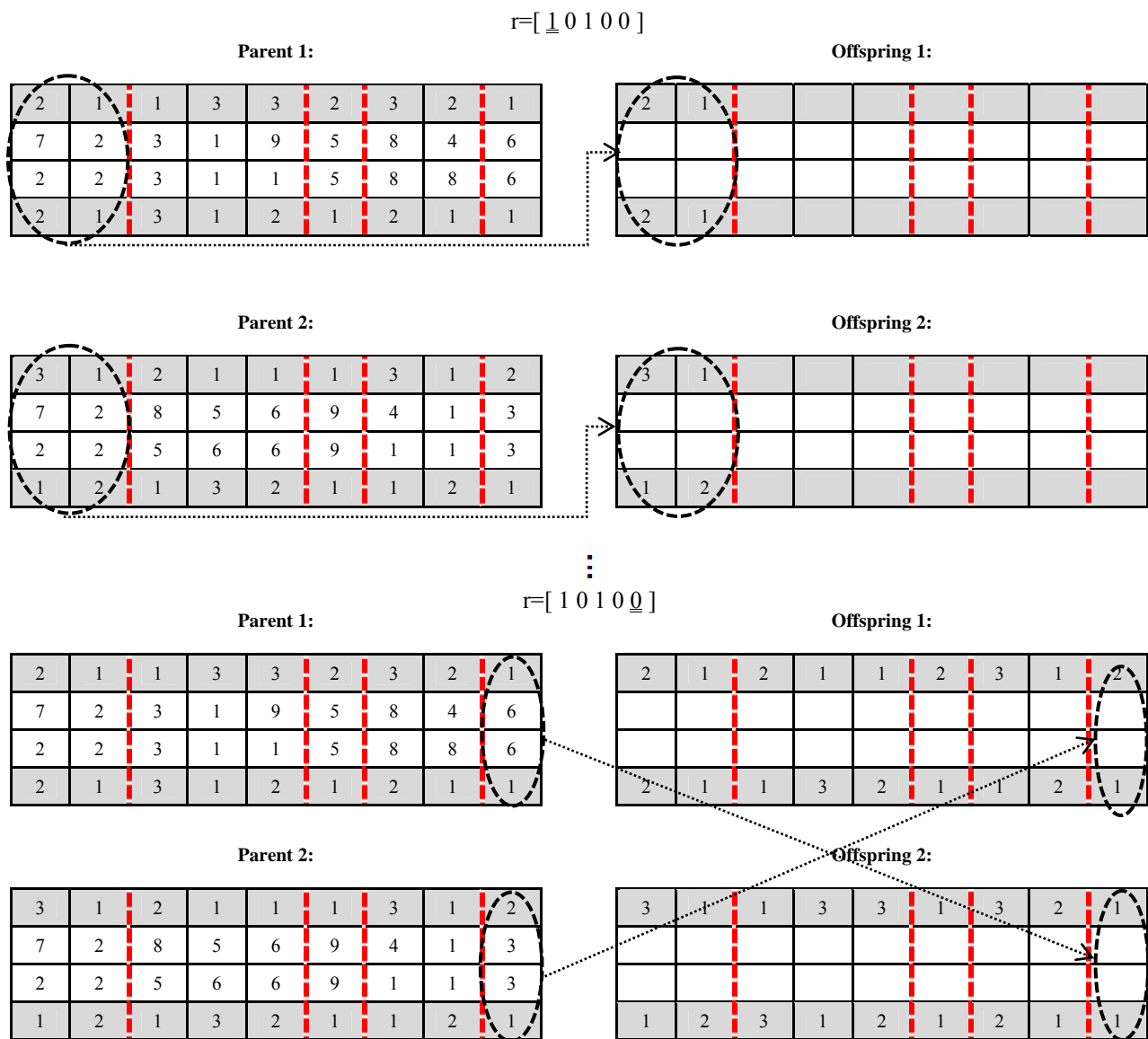


Fig 3. Proposed 'customer transfer' method for crossover operator for machine & priority orders rows of sample chromosome

Applying first step may improve the tardiness costs. Second step may decrease the processing time because the resulted earlier start time cause to shorter processing time (processing time is linear function of start time) and may decrease tardiness costs. Third step may decrease machine deteriorating costs and tardiness costs because it avoids overload on one machine that lead to increased machine deteriorating costs and tardiness costs. It is obvious that presence of many jobs on one machine will increase the delay time of jobs to enter processing and so tardiness costs. Pseudo-code of Proposed Heuristic Algorithm is presented in Table 3. In order to assign the jobs uniformly to machines, i.e. applying step 3 we define *expected* and *surplus* measures. *Expected* is quotient of total number of jobs divided by number of machines and surplus=N-

$m \times \text{Expected}$, i.e. *expected* determines the possible uniform assignment of jobs to machines and *surplus* calculates the number of additional jobs according to amount of *expected* e.g., if total number of jobs is 23 and number of machines is

$$4 \text{ then } \text{expected} = \frac{23}{4} = 5.75 \text{ and } \text{surplus} = 23 - 4 \times 5 = 3.$$

Moreover, *target* measure is defined as a binary vector with length of number of machines $\times 1$ in which 0 means that expected number of jobs have assigned to corresponding machine and 1 means that one job besides the expected number of jobs has assigned to it. Finally, *machine* vector represents the assignment of jobs to machines. Fig 6. illustrates above-mentioned calculations for an example solution after applying step 3 of Heuristic Algorithm.

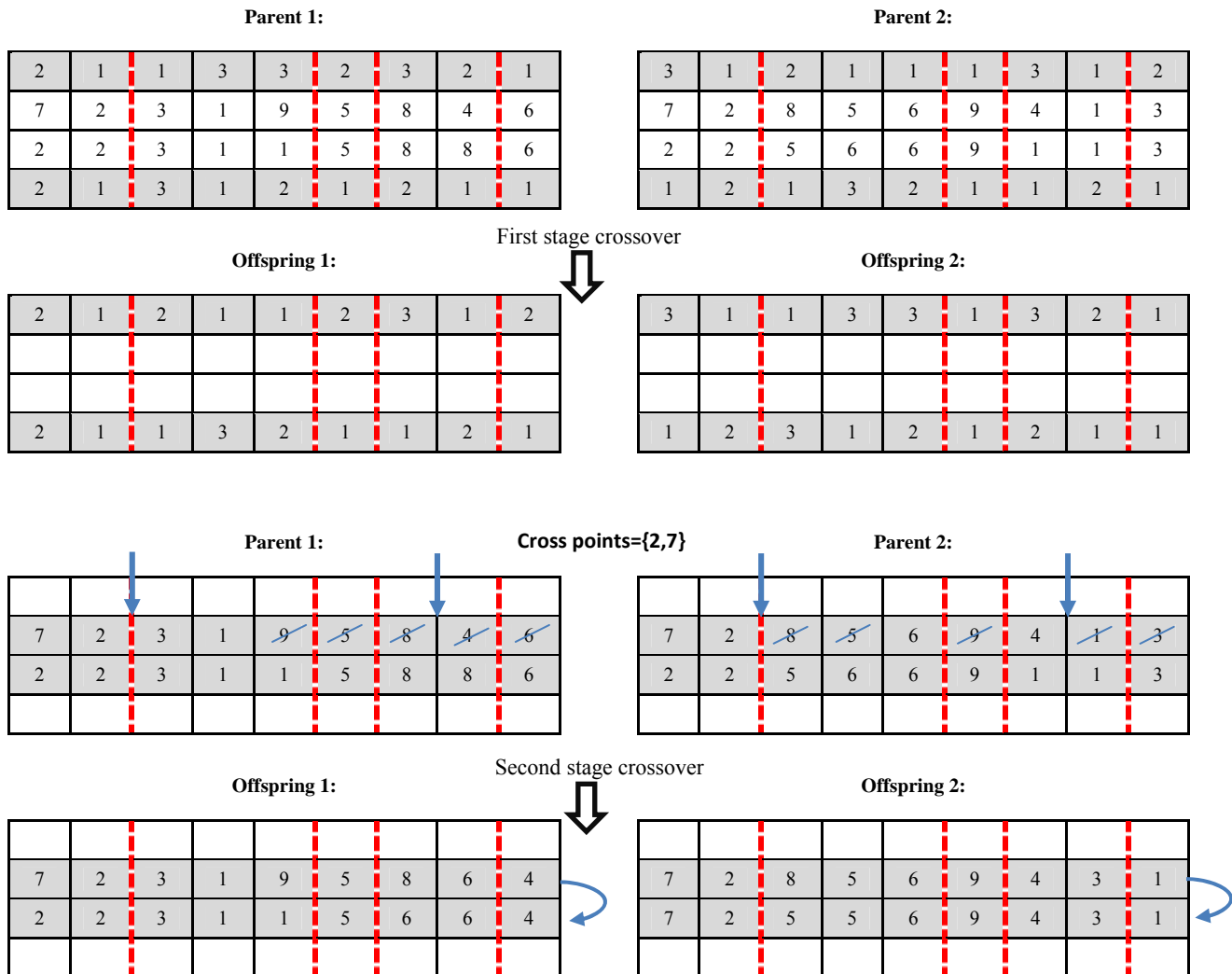


Fig 4. Proposed crossover operator for sample chromosomes

3.6. Details of the hybrid genetic algorithm

3.6.1. Input parameters

Input parameters for the proposed Hybrid Genetic Algorithm (HGA) are as follows:

Population size=80(HGA), 100(GA)

Max generation=100(HGA), 200(GA)

Probability of crossover=0.50

Probability of mutation=0.25

Probability of random solutions in each generation=0.05

Probability of elitist solutions selection=0.20

Pseudo-code of proposed Hybrid Genetic Algorithm is presented in Table 4.

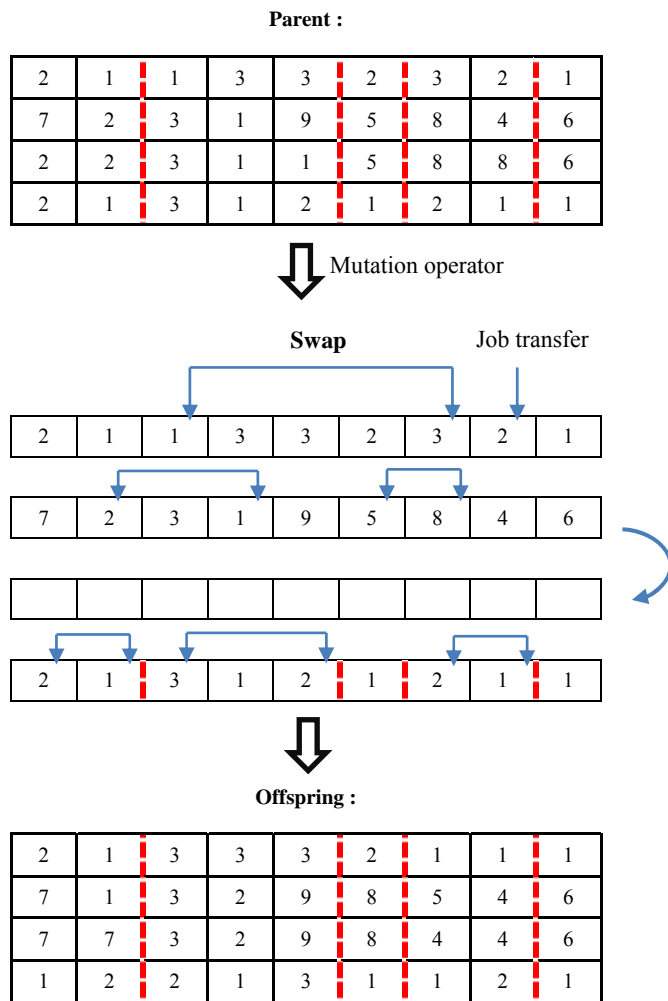


Fig 5. Proposed mutation operator for sample chromosome

4. Computational Results

In this section, we evaluate the performance of the suggested hybrid GA and traditional GA in comparison with the proposed MILP model. The proposed approaches are implemented in three cases (i.e., small, medium and large size test problems). It should be noted that according to the batch delivery scheduling problem literature, even in single machine case, excluding the deteriorating jobs and machines, CPLEX is only capable of solving small size problems (e.g., (Ahmadizar & Farhadi, 2015)). Consequently, the size of the test problems are defined regarding the high complexity of the problem.

The proposed mathematical model is coded in the GAMS 24 optimization software which uses the CPLEX solver and all the empirical experiments are run on a Intel® core™ i5 2.67GHz processor with 4 GB memory RAM. Also, the time limitation of 7200 seconds are considered for each run of CPLEX. The implementation results are given in Table 5. From the results of Table 5 we can observe that in small size test problem CPLEX have a good performance, but, although CPLEX is one the most powerful commercial solvers for solving optimization models, because of the high complexity of the MILP model regarding the number of constraints, it is inefficient in when the size of the problem increases. Also, to evaluate the results, runtime of each method, as well as its error are reported. In order to calculate the amount of error for proposed metaheuristics, each algorithm is executed 25 times with different random initializations and results are recorded. Amount of error for GA and HGA algorithm is calculated by Eq. (20) in which sol_i is the solution of algorithm in the i th implementation and optimum is the best solution of algorithm among 30 different outcomes.

$$Error = \frac{\sum_{i=1}^{30} |sol_i - optimum|}{30} \quad (20)$$

As can be seen from Table 5, in small and medium size problems, two metaheuristics have similar performances but in the case of large size problem hybrid GA reaches the global solution in the less time and error in comparison with traditional genetic algorithm. As seen in this table, presence of multiple machines improves the amount of total costs in comparison with single machine, moreover hybrid genetic algorithm outperforms traditional genetic algorithm.

Table 3
Pseudo-code of proposed Heuristic algorithm

```

Inputs: a solution (a chromosome), C (orders vector), N (total number of jobs), m (number of machines),
Outputs: improved solution
//Step 1
Set values of "order priority row" cells of each customer section to a permutation between 1 and  $O_i$  based on EDD (earliest due date) rule ( $O_i$  is number of  $i$ th customer's orders/jobs)
//Step 2
Sort and unique values of "batch number row" and set as b-set (identify used batch sets)
For i ← 1 to number of b-set (for each batch)
Identify the corresponding jobs
For j ← 1 to number of corresponding jobs -1
If values of genes 'j' and 'j+1' in the "machine row" are equal Then
Assign job 'j+1' to another machine
End If
EndFor
EndFor
// Step 3
Calculate Expected =  $\frac{N}{m}$  (Expected is the number of jobs that uniformly assign to machines, e.g., if N=23, m=4 so expected=5)
Calculate surplus = N - m * Expected (Remainder is the additional jobs, e.g., if N=23, m=4 so expected=5 and surplus=3)
Calculate Target = [1 1 ... 0 0]. (The number of ones in the Target matrix = surplus. Number of zeroes in Target vector = m - surplus. 0 means that the corresponding machine has Expected number of jobs and 1 means the machine has expected+1 number of jobs)
Calculate Machine = [Expected+1 Expected+1 ... Expected Expected] (assignment of jobs to machines)
Set jobs to machines based on Machine vector
//end of Heuristic Algorithm
    
```

Table 4
Pseudo-code of proposed hybrid genetic algorithm

```

Inputs: C (orders vector), N (total number of jobs), m (number of machines), PopSize (population size), Max-generation (number of generations), Pc (probability of crossover), Pm (probability of mutation), Prandom (probability of random solutions in each population), Pelitist (probability of selecting elitist solutions in each population)
Outputs: the best solution, minimum total costs and optimized assignment of jobs to machines
//initialization
Create Popsiz chromosomes
For each chromosome
Set values of "machine row" of the chromosome to random numbers between 1 and m
Set values of "initial batches row" of the chromosome to random permutations from 1 to N
Set values of "batch number row" cells of each customer section to random amounts selected from corresponding cells of "initial batches row"
Set values of "order priority row" cells of each customer section to random permutations between 1 and  $O_i$ 
Measure fitness of constructed solution based on input data set
//iterations
For I from 1 to max generation
//selection
Sort the individuals based on their fitness
Select ((1 - pc - Pm - Prandom) * Popsiz) top individuals for next generation
Generate (Prandom * Popsiz) random individuals for next generation
Select (pc * popsiz) individuals base on roulette wheel for crossover
//reproduction
For each pair in (Pc * Popsiz) selected individuals
Conduct the crossover operator to generate two off springs and select for next generation (pc * popsiz) of them
Select (Pm * Popsiz) individuals of off springs for mutation mutate each individual selected for mutation
Apply heuristic algorithm on mutated individuals and transfer them to next generation
//end of iteration
Select the fittest individual as the output of algorithm
//end of genetic algorithm
    
```

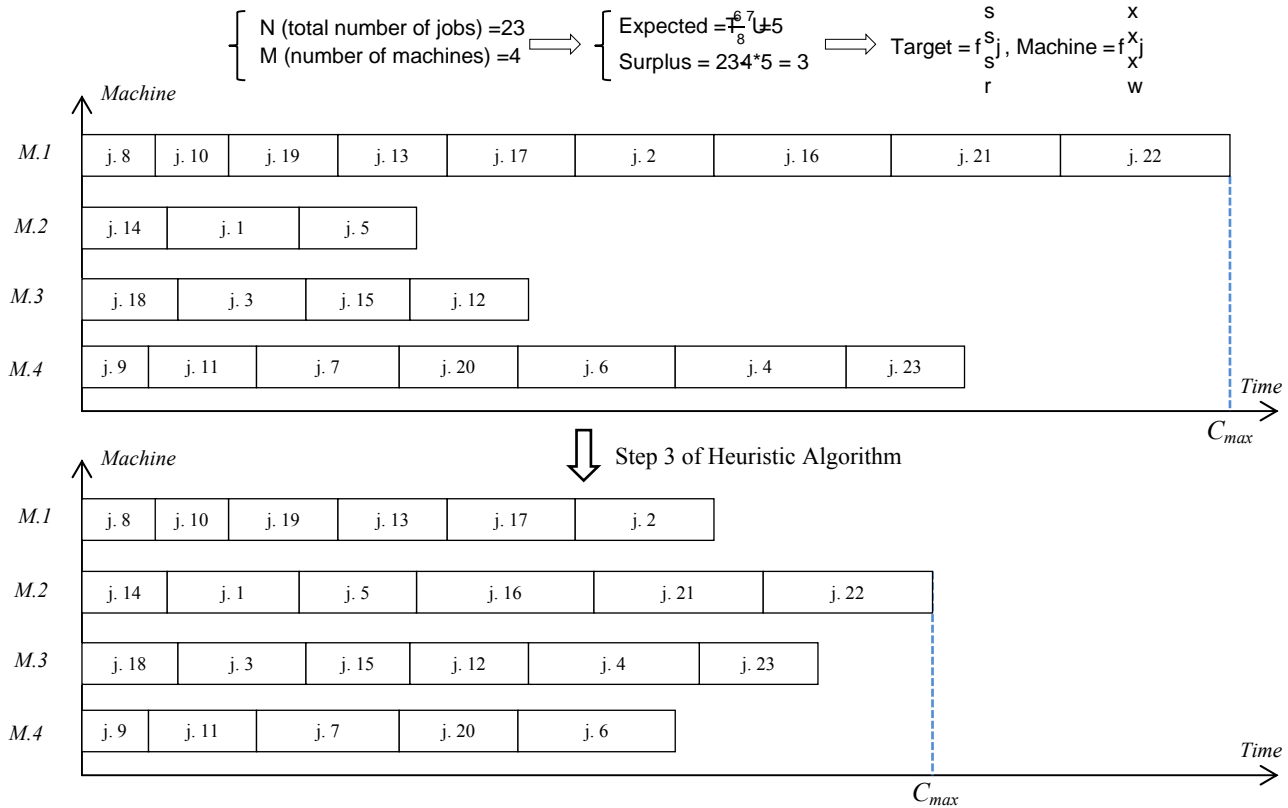


Fig. 6. Applying step 3 of proposed heuristic algorithm for example solution.

Table 5
Computational results of proposed approaches on test problems

Test problems:	Solution Results			CPLEX		GA		Hybrid GA	
	Objective function	Sequence on machines	batches	Run Time (s)	Error	Run Time (s)	Error	Run Time (s)	Error
<i>Small size:</i>									
N=4,m=1, C=[2 2]	1094.40	M1:2.2→2.1*→1.2→1.1	B1:1.1, B2:2.1 B3:1.2, B4:1.1	1.42	0	2.12	0	2.83	0
N=4,m=1, C=[2 2 2]	3845.70	M1: 2.2→2.3→2.1→1.3→ 1.2→1.1	B1:2.2, B2:2.3 B3:2.1, B4:1.3 B5:1.2, B6:1.1	13.128	0	3.78	0	3.69	0
<i>Medium size:</i>									
N=4,m=1, C=[2 2 2 2]	9635.30	M1:1.4→2.2→2.3→2.1 1→2.4→1.3→1.2→1.1	B1:1.4, B2:2.2 B3:2.3, B4:2.1 B5:2.4, B6:1.3 B7:1.2, B8:1.1	2760	0	5.31	0	4.63	0
N=8,m=1, C=[4 4]	4712.90	M1:2.2→1.1→4.1→2.1 1→3.1→4.2→3.2→1.2	B1:2.2, B2: B3:1.1,4.1, B4:2.1 B5:3.1, B6:4.2 B7:3.2, B8:1.2	2871	0	5.12	0	4.62	0
<i>Large size:</i>									
N=9,m=3, C=[2 3 1 2 1]	588.9	M1:1→4→9 M2:2→5→7 M3:6→3→8	B1:1.3, B2:1.1,2.1 B5: 2.2,3.2 B6:1.2, B7:1.5 B8:2.4, B9:1.4	-	-	10.75	%3	7.33	0
N=10,m=3, C=[2 3 1 2 1 1]	747.91	M1:1→4→9 M2:5→2→7→10 M3:6→3→8	B1:1.3, B4:2.2,3.2 B5:1.1,2.1, B6: 1.4 B7:1.2, B8:1.5 B9:1.6, B10:2.4	-	-	16.39	%6	9.08	%3

* 2.1: second job/order of first customer

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