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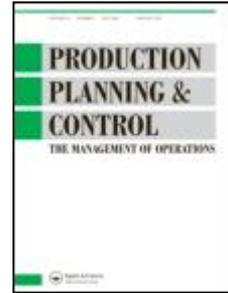
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A mathematical model and genetic algorithm-based approach for parallel two-sided assembly line balancing problem

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Abstract: Assembly lines are usually constructed as the last stage of the entire production system and efficiency of an assembly line is one of the most important factors which affect the performance of a complex production system. The main purpose of this paper is to mathematically formulate and to provide an insight for modelling the parallel two-sided assembly line balancing problem, where two or more two-sided assembly lines are constructed in parallel to each other. We also propose a new genetic algorithm (GA)-based approach in alternatively to the existing only solution approach in the literature, which is a tabu search algorithm. To the best of our knowledge, this is the first formal presentation of the problem as well as the proposed algorithm is the first attempt to solve the problem with a GA-based approach in the literature. The proposed approach is illustrated with an example to explain the procedures of the algorithm. Test problems are solved and promising results are obtained. Statistical tests are designed to analyse the advantage of line parallelisation in two-sided assembly lines through obtained test results. The response of the overall system to the changes in the cycle times of the parallel lines is also analysed through test problems for the first time in the literature.

Keywords: parallel two-sided assembly lines; assembly line balancing; production planning; genetic algorithm; meta-heuristics; artificial intelligence

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A Mathematical Model and Genetic Algorithm based Approach for Parallel Two-Sided Assembly Line Balancing Problem

Assembly lines are usually constructed as the last stage of the entire production system and efficiency of an assembly line is one of the most important factors which affect the performance of a complex production system. The main purpose of this paper is to mathematically formulate and to provide an insight for modelling the parallel two-sided assembly line balancing problem, where two or more two-sided assembly lines are constructed in parallel to each other. We also propose a new genetic algorithm based approach in alternatively to the existing only solution approach in the literature, which is a tabu search algorithm. To the best of our knowledge, this is the first formal presentation of the problem as well as the proposed algorithm is the first attempt to solve the problem with a genetic algorithm based approach in the literature. The proposed approach is illustrated with an example to explain the procedures of the algorithm. Test problems are solved and promising results are obtained. Statistical tests are designed to analyse the advantage of line parallelisation in two-sided assembly lines through obtained test results. The response of the overall system to the changes in the cycle times of the parallel lines is also analysed through test problems for the first time in the literature.

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1. Introduction

Assembly lines have been utilised successfully to produce large-volume high-quality standardised homogeneous products, and have been of interest for both academia and industry for decades. An assembly line is a sequential organisation of workstations (or operators) linked by a conveyor belt or material handling system on which semi-finished products are moved from one workstation to another (Ozdemir and Ayag 2011). Parts are added on the moving semi-finished products in sequence until the final assembly is produced. A group of tasks is performed in each workstation by considering

capacity of the workstation and precedence relationships among tasks, where precedence relationships are usually caused by the technological requirements or organisational structures (Tonelli *et al.* 2013). Sum of processing times of all tasks assigned to a workstation constitutes its workload time and workload time of a workstation cannot exceed the designated cycle time. Assembly line balancing problem is to determine which task will be accomplished in which workstation by assigning tasks to an ordered sequence of workstations considering aforementioned constraints (*i.e.* capacity constraints, assignment constraints, precedence relationships constraints, *etc.*). Tasks cannot be split into two or more pieces and each task must be assigned to exactly one workstation (Kucukkoc *et al.* 2013).

In accordance with the utilisation of operation sides, assembly lines can be classified as one-sided assembly lines and two-sided assembly lines. Tasks are performed on both left and right sides of the line in a two-sided assembly line system while only left or right side of the line is used in a one-sided assembly line system. Two-sided assembly lines are usually utilised to produce high-volume large-sized products, such as trucks and buses (Kucukkoc and Zhang 2014b). Some heuristic approaches were proposed by Lee *et al.* (2001), Hu *et al.* (2008), Ozcan and Toklu (2010) and Yegul *et al.* (2010); and some exact solution approaches were developed by Wu *et al.* (2008) and Hu *et al.* (2010) since the two-sided assembly line balancing problem was first introduced by Bartholdi (1993). Meta-heuristics have also been presented by Baykasoglu and Dereli (2008), Simaria and Vilarinho (2009), Ozcan and Toklu (2009), Ozbakir and Tapkan (2010), Ozcan (2010), Ozbakir and Tapkan (2011), Chutima and Chimklai (2012), and Khorasanian *et al.* (2013). Among these meta-heuristics, studies by Kim *et al.* (2000), Kim *et al.* (2009), Taha *et al.* (2011), Rabbani *et al.* (2012) and Purnomo *et al.* (2013) employed Genetic Algorithm (GA) based approaches to balance two-sided lines. As can

be comprehended from these studies, there exist numerous successfully implemented GA based approaches in the literature for two-sided assembly line balancing problems.

There is another type of line configuration called parallel assembly line system, where two or more lines are located in parallel to each other to maximise the use of shared resources and tools. The idea of balancing more than one assembly line with a common set of resources was first introduced by Gökçen *et al.* (2006). Gökçen *et al.* (2006) proposed new procedures and a mathematical model on the single model assembly line balancing problem with parallel lines. Few researchers followed Gökçen *et al.* (2006) and a novel ant colony optimisation based algorithm was proposed by Baykasoglu *et al.* (2009) for the *Parallel Assembly Line Balancing Problem (PALBP)*. Cercioglu *et al.* (2009) proposed a simulated annealing approach to solve PALBP and compared results obtained from the algorithm with the results of existing heuristic algorithm proposed by Gökçen *et al.* (2006). The first multi-objective tabu search algorithm is presented for PALBP by Ozcan *et al.* (2009) and its performance was tested on a set of well-known problems in the literature. Another mathematical model of the PALBP was developed by Scholl and Boysen (2009) along with an exact solution procedure. Kara *et al.* (2010) suggested a fuzzy goal programming model that can be used for balancing parallel assembly lines. Ozcan *et al.* (2010a) addressed parallel mixed-model assembly line balancing and sequencing problem with a simulated annealing approach to maximise the line efficiency by ensuring smooth workload distribution among workstations. Ozbakir *et al.* (2011) developed a multiple-colony ant algorithm for balancing bi-objective parallel assembly lines while Kucukkoc and Zhang (2014a, 2014b) considered the model sequencing problem as well as the line balancing problem on mixed-model parallel two-sided assembly lines and proposed agent based ant colony optimisation solution approaches. Please refer to Lusa (2008) and Zhang and

Kucukkoc (2013) for a detailed survey on multiple and parallel assembly line balancing problems.

The combination of above mentioned configurations, namely *parallel two-sided assembly lines*, are also frequently constructed in industry in production of large-sized items. Although vast numbers of researches have been carried out on traditional configurations of assembly lines in the literature, there is only one research concerning *Parallel Two-Sided Assembly Line Balancing Problem (PTALBP)*. The concept of parallel two-sided assembly lines was first described by Ozcan *et al.* (2010b). Ozcan *et al.* (2010b) introduced and defined the PTALBP and proposed a tabu search algorithm to solve the combined well-known test problems in the literature. Obtained results were compared with theoretical minimum number of workstations to show the performance of the proposed algorithm. The research also demonstrated that parallelisation of two-sided lines helps lower total number of workstations, but no statistical technique was used for this aim.

GAs as well as other evolutionary approaches have been applied to line balancing problems earlier with success. There is continuing work in applying GA for various types of line balancing problems, *i.e.* Leu *et al.* (1994), Rubinovitz (1995), Kim *et al.* (2000), Rekiek *et al.* (2001), Goncalves (2002), Simaria and Vilarinho (2004), Zhang *et al.* (2005), Haq *et al.* (2006), Levitin *et al.* (2006), Suwannarongsri *et al.* (2007), Zhang *et al.* (2008), Hwang and Katayama (2009), Yu and Yin (2010), Chica *et al.* (2011), Akpinar and Bayhan (2011), and Kucukkoc *et al.* (2013). In particular, Kim *et al.* (2000), Kim *et al.* (2009), Taha *et al.* (2011), Rabbani *et al.* (2012), and Purnomo *et al.* (2013) have solved two-sided assembly line balancing problems with different GA based approaches but none of them have considered line parallelisation as an additional specification. On the contrary of its successful implementations on various line systems,

PTALBP has not been addressed using any GA based technique. Therefore, there is neither GA based nor evolutionary approach published concerning parallel two-sided assembly lines. This is the main motivation of why a GA based approach is proposed in solving the addressed problem in this research.

Evidence of the need for this research is showed by the lack of literature on developing the mathematical model of the PTALBP and presenting the positive effect of line parallelisation on two-sided lines, statistically. The need for this research is also guided by the gap in the literature on balancing more than one two-sided assembly line with a common set of resources using an evolutionary based approach, such as GA. Moreover, the response of the whole line system against the changes in the cycle times of the parallel lines is demonstrated for the first time in the literature. From the managerial point of view, among the available solutions, line managers can easily pick up a solution for a specific combination of cycle times of the parallel lines. This helps them make decision especially when there is change in model demands.

The rest of the paper is organised as follows. Section 2 presents the main characteristics of the problem along with the mathematical model and assumptions considered. The proposed approach is described in Section 3 and illustrated with an example in Section 4. Section 5 reports and statistically analyses the results of the computational study. Section 6 concludes with the findings of the research and the future research directions. Some graphs related to the computational study are also depicted in the Appendices.

2. Problem Statement

2.1. Main characteristics

To maximise the use of shared tools and minimise idle times of the entire production

system, two or more two-sided lines – on which two or more similar product models that have similar production processes are produced – are located in parallel to each other. Such a configuration is called parallel two-sided assembly line system and can be typically illustrated as in Figure 1. The parallel two-sided assembly line balancing problem is balancing two or more two-sided assembly lines, which are constructed in parallel to each other. A common set of resources is shared among the lines and the main objective is allocating tasks to the workstations to optimise a performance measure (*i.e.* number of workstations, line efficiency, *etc.*) by considering technological priorities, capacity constraints and some other possible constraints caused by organisational structures or technological requirements. Different product models are produced on each of the two-sided assembly lines, represented by h (where $h = 1, \dots, H$); and each product model has its own set of tasks ($i = 1, \dots, n_h$). These tasks are performed according to the known precedence relationships among tasks. P_h represents set of precedence relationships on line h , where $(r, s) \in P_h$ represents that Task- r must be completed to be able to assign Task- s . Each task, which is performed on line h , needs a certain amount of processing time symbolised with t_{hi} ; and each line consists of a series of workstations ($k = 1, \dots, K$) (Ozcan *et al.* 2010b).

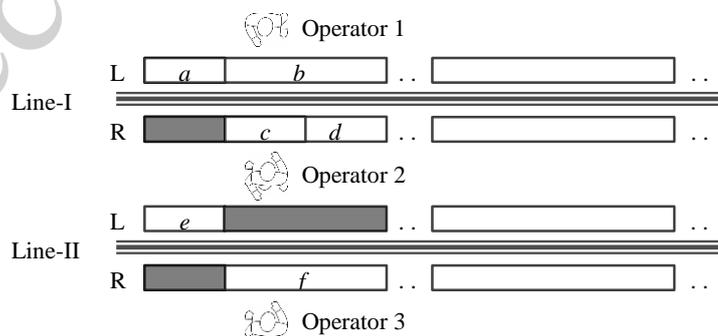


Figure 1. Typical illustration of parallel two-sided assembly lines, adapted from (Ozcan *et al.* 2010b).

The main advantage of parallel two-sided assembly lines is the flexibility of utilising

multi-line stations between two adjacent lines. Operators located in stations between two adjacent lines can perform tasks from both lines. By this way, idle times are reduced and system utilisation is increased. As can be seen in Figure 1, three operators are needed to perform Tasks-*a* – Task-*f*, and Operator-2 first completes Task-*e* at the left side station of Line-II, and then Task-*c* and Task-*d* at the right side station of Line-I. Please note that the shades in the figure symbolise idle times (Ozcan *et al.* 2010b).

It should be noted that more attention is needed when balancing two-sided assembly lines because tasks, which have precedence relationships with each other and are performed on different sides of the lines, must be assigned considering finishing time of previously assigned tasks. Let P_1 denotes the set of precedence relationships of Line-I. If $(a, b) \in P_1$ and $(a, c) \in P_1$, then Task-*b* and Task-*c* can be initialised after completion of Task-*a*, which may be performed at the other side of the line. This phenomenon is called *interference* in the literature and the violation of this rule yields infeasible balancing solutions.

Another significant advantage of this line system is that each line may have a different cycle time (C_h), which means that each line may have a different throughput rate contributing to the flexibility. When two lines which have different cycle times are subject to balancing, a common cycle time should be used to assign tasks in each cycle. Gökçen *et al.* (2006) used least common multiple (*LCM*) based approach for different cycle time situation of two parallel lines (Ozcan *et al.* 2010b). In this approach (Gökçen *et al.* 2006):

- Least common multiple of the cycle times is found.
- Line divisors (ld_1 and ld_2) are calculated through dividing the *LCM* value by the cycle times of Line-I and Line-II (C_1 and C_2), respectively.
- Task times of the product models produced on the Line-I and Line-II are multiplied by ld_1 and ld_2 , separately.

- *LCM* is determined as the common cycle time (*C*) of the lines and the lines are balanced together.

To characterise the PTALBP more clearly and provide an insight for modelling of PTALBP and utilisation of multi-line stations, a numerical example is given below. Data given in Table 1 is used as input for the example problem and a possible balancing solution for the considered problem is exhibited in Figure 2 under 12 time-units cycle time constraint for both of the lines.

Table 1. Data for numerical example.

Task No	Line-I			Line-II		
	Side	Processing Time	Immediate Predecessors	Side	Processing Time	Immediate Predecessors
1	Left	2	-	Left	2	-
2	Either	4	1	Right	2	-
3	Right	3	-	Left	4	-
4	Either	3	2	Either	1	2
5	Left	6	1	Right	3	2
6	Either	4	5	Either	3	1
7	Left	5	4,6	Left	5	3,4,6
8	Either	1	4	Either	4	5
9	Either	3	8	Left	4	7
10	Right	2	3	Either	3	7,8
11	Either	2	10	Either	3	-
12	Left	4	11	Left	2	9
13	Either	3	7	Right	5	10
14	Left	4	13	Either	3	11
15	Either	2	12	Left	6	12
16	Either	3	14,15	Either	3	13,15
17	Left	2	16	Either	6	14
18	-	-	-	Left	2	16
Total Time		53			61	

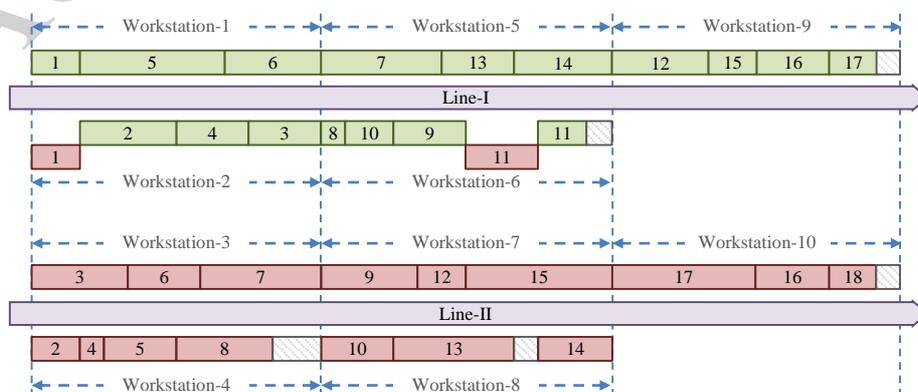


Figure 2. A possible balancing solution using multi-line stations.

Figure 2 shows the utilisation of multi-line stations between the adjacent two-sided lines located in parallel to each other. Numbers inside bars denote task numbers while lengths of the bars correspond to processing times of tasks. Dashed areas represent unavoidable idle times caused by capacity and precedence relationships constraints. As it could be seen from the figure, a total of 10 workstations are needed to perform a total of 35 tasks on both of the lines. With the construction of multi-line stations, operator located in workstation-2 on Line-I performs Task-1 from Line-II first, followed by Task-2, Task-4, and Task-3 belonging to Line-I. Similarly, Task-11 from Line-II is completed by workstation-6 located on Line-I. Thus, operators located in workstation-2 and workstation-6 on Line-I contribute to performing tasks on Line-II as well as their main job on Line-I. It should be noted here that Task-14 on Line-II cannot be initialised unless its predecessor task (Task-11) is completed by workstation-6 on the other side of the line. This is one of the most challenging issues in solving PTALBPs and that is why unavoidable idle time occurs before Task-14 on Line-II.

If the lines were balanced individually (without multi-line stations), theoretical minimum number of workstations for Line-I and Line-II could be calculated as $minK_1 = [53/12]^+ = 5$ and $minK_2 = [61/12]^+ = 6$, respectively; simply using the well-known formula $minK_h = [Total\ Task\ Time/Cycle\ Time]^+$, where $[X]^+$ denotes the smallest integer greater than or equals to X . Whereas theoretical minimum number of workstations ($minK$) decreases to 10 ($minK = [114/12]^+$), one lower than the sum of independent balancing solutions ($5 + 6 = 11$), with the opportunity of assigning tasks into a more diversified positions thanks to multi-line stations when the lines are balanced together.

2.2. Mathematical model

The notation used in the study can be summarised as below to describe the problem:

2.2.1. Indices:

h : line index ($h = 1, \dots, H$), where H represents total number of lines,

i : task index ($i = 1, \dots, n_h$), where n_h represents total number of tasks on line h ,

j : side of the line, $j = \begin{cases} 0 & \text{indicates left side} \\ 1 & \text{indicates right side} \end{cases}$,

k : station index ($k = 1, \dots, K$), where K represents total number of utilised workstations.

2.2.2. Decision Variables:

$$X_{hijk} = \begin{cases} 1 & \text{if task } i \text{ is assigned to workstation } k, \text{ on side } j \text{ of line } h \\ 0 & \text{otherwise} \end{cases}.$$

2.2.3. Parameters:

t_{hi} : Processing time of task i on line h ,

P_h : Set of precedence relationships in precedence diagram of line h ,

C : Common cycle time of the lines,

ZP : Set of pairs of tasks that must be assigned to the same workstation, *positive zoning*,

NP : Set of pairs of tasks that cannot be assigned to the same workstation, *negative zoning*.

2.2.4. Indicator Variables:

t_{hi}^s : Starting time of task i on line h ,

q_k : Queue number that station k is utilised on,

$$S_k = \begin{cases} 0 & \text{if station } k \text{ is utilised on left side of the first line} \\ 1 & \text{otherwise} \end{cases},$$

$$z_k = \begin{cases} 1 & \text{if station } k \text{ is utilised} \\ 0 & \text{otherwise} \end{cases},$$

$$U_{hjk} = \begin{cases} 1 & \text{if station } k \text{ is utilised on side } j \text{ of line } h \\ 0 & \text{otherwise} \end{cases},$$

σ : Variable, ($\sigma = h + 1, \dots, H$),

β : Variable, ($\beta \in \{0, 1\}$),

$$c = \begin{cases} 1 & \text{if } j = 1 \text{ and } \beta = 1 \\ 0 & \text{otherwise} \end{cases},$$

$$\mu = \begin{cases} 1 & \text{if } (\sigma - h) = 1 \text{ and } \beta = 1 \\ 0 & \text{otherwise} \end{cases},$$

$$R_{hrs} = \begin{cases} 1 & \text{if tasks } r \text{ and } s \text{ are assigned to the same workstation on line } h \\ 0 & \text{otherwise} \end{cases}.$$

2.2.5. Objective Function:

The objective function used in this study is obtained from the modification of objective functions used in the previous studies (see Chiang (1998) and Ozcan *et al.* (2010b)) and is presented in Equation (1). This nonlinear objective function represents sum of squares of each workstation's workload. So, maximising this objective function helps to reduce the number of stations.

$$\text{Max } Z = \sum_{h=1}^H \sum_{j \in \{0,1\}} \sum_{k=1}^K \left(\sum_{i=1}^{n_h} t_{hi} X_{hijk} \right)^2 \quad (1)$$

2.2.6. Constraints:

$$\sum_{k=1}^K X_{hijk} = 1, \quad \forall i \in n_h; \quad \forall h \in H; \quad \forall j \in \{0, 1\}. \quad (2)$$

$$\sum_{i=1}^{n_h} (t_{hi} + t_{hi}^s) X_{hijk} + S_k \left(\sum_{i=1}^{n_h} (t_{(h+1)i} + t_{(h+1)i}^s) X_{(h+1)i(j-1)k} \right) \leq Cz_k, \quad \forall k \in K; \quad \forall h = 1, \dots, H - 1; \quad \forall j \in \{0, 1\}. \quad (3)$$

$$\sum_{i=1}^{n_h} X_{hijk} - n_h U_{hjk} \leq 0, \quad \forall k \in K; \quad \forall h \in H; \quad \forall j \in \{0, 1\}. \quad (4)$$

$$|j - 1|(U_{h\beta k} + U_{(\sigma-\mu)jk}) + j(U_{h(j-1+c)k} + U_{\sigma jk}) = 1, \quad \forall k \in K; \quad \forall h \in H; \quad \forall j \in \{0, 1\}; \quad \forall \sigma = h + 1, \dots, H; \quad \forall \beta \in \{0, 1\}. \quad (5)$$

$$\sum_{k=1}^K q_k(X_{hrjk} - X_{hsjk}) + R_{hrs}(t_{hr}^s + t_{hr} - t_{hs}^s) \leq 0, \quad \forall h \in H; \quad \forall j \in \{0, 1\}; \quad \forall (r, s) \in P_h. \quad (6)$$

$$\sum_{k=1}^K X_{hajak} - \sum_{k=1}^K X_{hbjk} = 0, \quad \forall (a, b) \in ZP; \quad \forall h \in H; \quad \forall j \in \{0, 1\}. \quad (7)$$

$$X_{hajak} + X_{hbjk} \leq 1, \quad \forall (a, b) \in ZN; \quad \forall h \in H; \quad \forall j \in \{0, 1\}; \quad \forall k \in K. \quad (8)$$

The main objective of the model given in Equation (1) is to minimise the number of workstations by maximising sum of squares of each workstation's workload. Constraint (2) ensures that all tasks are assigned to a station and each task is assigned only once. Constraint (3) represents cycle time constraint that assures each task is executed before the cycle time. Constraints (4) and (5) ensure that an operator working at station k can perform additional task(s) from only one adjacent line; unless station k is not utilised on left side of the first line or on right side of the last line; *i.e.* if an operator is located on right side of the first line ($h = 1, j = 1$), that operator can perform additional tasks from only left side of the second line ($h = 2, j = 0$) as well as his/her main job. That operator cannot perform any job from left side of the first line ($h = 1, j = 0$), or right side of the second line ($h = 2, j = 1$), as it is not possible a direct communication with those tasks assigned to these stations. Please refer to Section 2.2.4 for explanations on indicator variables c and μ given in constraint 5. Constraint (6) ensures that the precedence relationships are not violated and completion times of tasks are considered to avoid interference. Given a task pair $(r, s) \in P_h$, where r is one of the predecessors of s , then s can be initialised after r is completed. Constraints (7) and (8) demonstrate

positive and negative zoning constraints, respectively. As noted above, ZP is the set of pairs of tasks that must be assigned to the same workstation, while ZN is the set of pairs of tasks that cannot be assigned to the same workstation.

2.3. Assumptions

The assumptions considered in the study are as follows:

- Only one product model is assembled on each line, so total number of lines equals to total number of product models.
- Each product model has its own precedence relationships diagram.
- The precedence relationships and task times of each product model are known.
- The operators have no preference about the tasks and workstations.
- Walking times of the operators are ignored.

3. Proposed GA based Approach for PTALBP

GA is an efficient random search algorithm originating from the evolutionary rules of the nature population. Its solution approach is motivated by the biological process of natural selection and the solution of an optimisation problem is encoded as chromosome where the specific parameters of solution (called genes) are located on the chromosome (Suresh *et al.* 1996). Each individual (chromosome) corresponds to a possible solution and its survival chance through generations is characterised by its fitness value, which is defined in accordance with the objective function. A finite set of individuals constitutes population and usually its size remains fixed through generations. The initial population is built at random and the population is updated by generating new individuals, which replace the old ones, in subsequent iterations. New individuals are created by means of genetic operators, crossover and mutation, and the iterations are terminated when the stopping criterion is satisfied (Borisovsky *et al.* 2013). The characteristics of the implemented GA approach within the scope of this study are explained below.

3.1. General outline

The outline of the proposed GA based algorithm is exhibited in Figure 3. As can be seen from the figure, the algorithm starts by generating an initial population, which consists of a predefined number (population size) of chromosomes, and continues with the evaluation of created chromosomes. Genetic operators (crossover and mutation) are performed and some completely new chromosomes are also generated randomly with the probability of 2% to keep diversity and avoid early convergence.

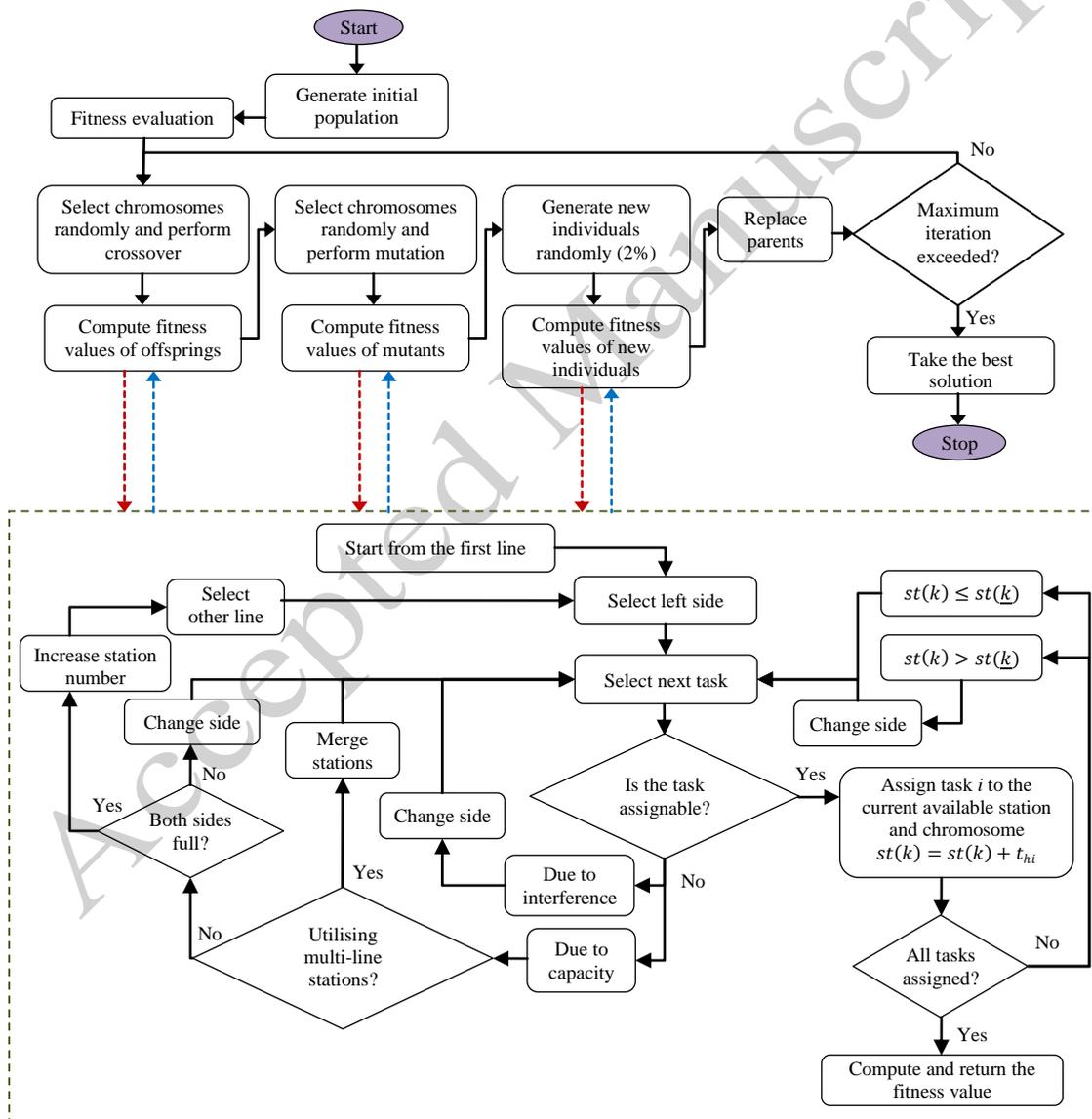


Figure 3. Flowchart of the proposed algorithm, adapted from Kucukkoc and Zhang (2013).

After the fitness evaluation of new individuals (resulting from genetic operators and random generation), insufficient chromosomes in the population are replaced with better ones (if any). In the fitness evaluation process, tasks are assigned to the minimum numbered workstations as far as possible. This loop continues until the iteration number is exceeded. Finally, the chromosome which gives the best fitness value is selected as the best solution of the problem.

3.2. Initial population

The chromosome is made up of several genes which represent tasks (by the tasks' index numbers) in a sequence. So, each gene of the chromosome is an integer representing a task number of a sequence of tasks to be assigned to the stations. Different solutions and fitness values are examined by changing the order of the genes on the chromosome. Figure 4 represents a sample of task based chromosome which is used in this study. The length of the chromosome is characterised by the total number of tasks belong to the models. If we assume two product models with 9 tasks and 8 tasks, respectively, gene numbers lower than or equal to 9 belong to the Product Model-I (produced on Line-I). The remaining tasks (Task-10 – Task-17) belong to the Product Model-II (produced on Line-II) in an incremental order, *i.e.* Task-10 and Task-13 symbolise Task-1 and Task-4 for Product Model-II.

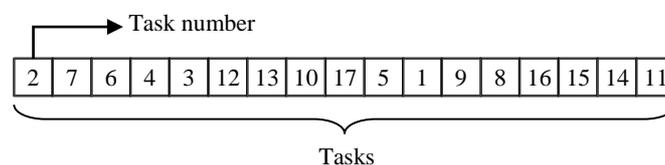


Figure 4. An example of task based chromosome representation.

Initial population is generated randomly using a heuristic algorithm, namely Comsoal (Arcus 1966), to start the GA. But first of all, tasks are grouped according to the line and preferred operation direction data. S1LE, S1RE, S2LE, and S2RE lists (named S

lists) are formed to separate tasks according to their input data. For example, S1LE list consists of tasks that can be assigned to left side of Line-I. After separating tasks, a Comsoal based heuristic procedure generates different chromosomes by selecting tasks from these lists until a population is obtained with a predetermined size. This procedure is exhibited in Figure 5.

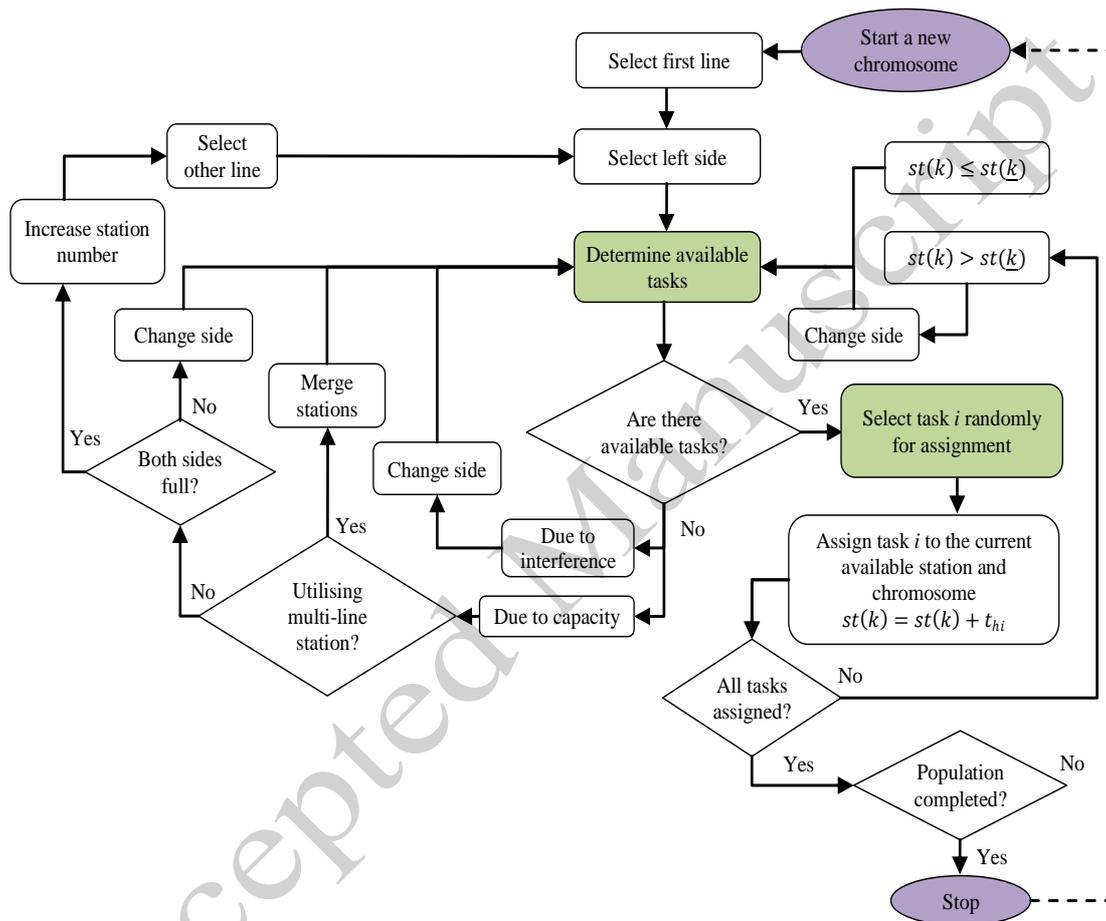


Figure 5. Generating initial population for the proposed algorithm, adapted from Kucukkoc and Zhang (2013).

The process flow as to how a chromosome is generated is also depicted in Figure 5; where available tasks mean those tasks which (i) satisfy capacity constraints of the current station, (ii) have no predecessors or all of their predecessor tasks are already completed and (iii) do not violate interference rule. For each side of each line, tasks with no predecessor and satisfy capacity constraints are selected randomly from relevant list and allocated to the chromosome one by one, then those tasks whose predecessors

have been processed and allocated to the chromosome, and so on.

Allocating tasks to the chromosome continues until all tasks are sequenced on the chromosome side by side and line by line. As only the tasks which have no predecessor or whose predecessors have been allocated are selected in each loop, infeasible task sequences are naturally filtered out during this process to prevent infeasible chromosomes (solutions) that violate precedence relationships. At this step, the length of each chromosome equals to the total number of tasks on both lines. As can be seen from the figure, if workload of the current station is larger than the workload of its mated station ($st(k) > st(\underline{k})$), then side is changed and candidate tasks for new side are considered. Afterwards, fitness values of the chromosomes are computed (decoding). During the task allocation process, if the current side of a line lies between two lines and there is no available task to be assigned from the current line but from the adjacent line, the multi-line station is utilised so that some tasks can be performed from the other line.

3.3. Decoding and fitness evaluation

Decoding procedure is processed by assigning tasks to workstations according to precedence relationships, unless cycle time is not exceeded. The sequence of tasks on the chromosome is considered while allocating tasks to the stations. The initial tasks on the chromosome are assigned in the earliest workstations as far as possible. If the next task in the sequence does not satisfy the precedence or capacity constraints, a new workstation is opened and the task is assigned to this workstation. Fitness value of each chromosome is computed when all tasks are assigned. Total number of workstations are also recorded for each chromosome in order to compare the obtained results with previous tabu search algorithm proposed by Ozcan *et al.* (2010b).

3.4. Selection, crossover and mutation

Crossover operator takes two parent individuals and produces two offsprings by combining and exchanging their elements. Roulette wheel (Baker 1987) is used to select parent chromosomes from the population to keep diversity and avoid local minima. Two-point crossover operator is applied to recombine the chromosomes. During this process, infeasible solutions are not allowed since missing parts of both offsprings are built according to precedence relationships. Selected two parents are divided into three sections: head, middle and tail. Cutting points, which cut each parent into three parts, are determined randomly for each parent pair. By this way, diversity is preserved in the population and it is enabled to search the solution space effectively. First offspring keeps the head and tail parts of the first parent and the middle part of the first offspring is filled by adding missing tasks according to the order in which they are contained in the second parent. Similarly, second offspring is formed by head part of the second parent, missing tasks according to the order in which they are contained in the first parent, and tail part of the second parent (Leu *et al.* 1994, Akpinar and Bayhan 2011). An example of the crossover procedure used in this study is given in Figure 6a. As could be seen from the figure, missing tasks of Offspring-1 are 5, 12, 15, 10, 11, 14, 18, 7, 8, and 6. These tasks appear in the sequence of 6, 5, 10, 15, 12, 11, 14, 8, 7, 18 on Parent-2, and constitute the middle part of Offspring-1.

Mutation is applied to add random changes to an individual and it plays a critical role in GA to keep diversity by changing the order of the genes dramatically. Roulette wheel selection strategy is applied so that an individual with a high fitness value will have more chance to be chosen as a parent than the ones with a lower fitness. To mutate a chromosome, two genes are selected randomly and swapped by considering precedence relationships among tasks. An example of mutation procedure is presented in Figure 6b.

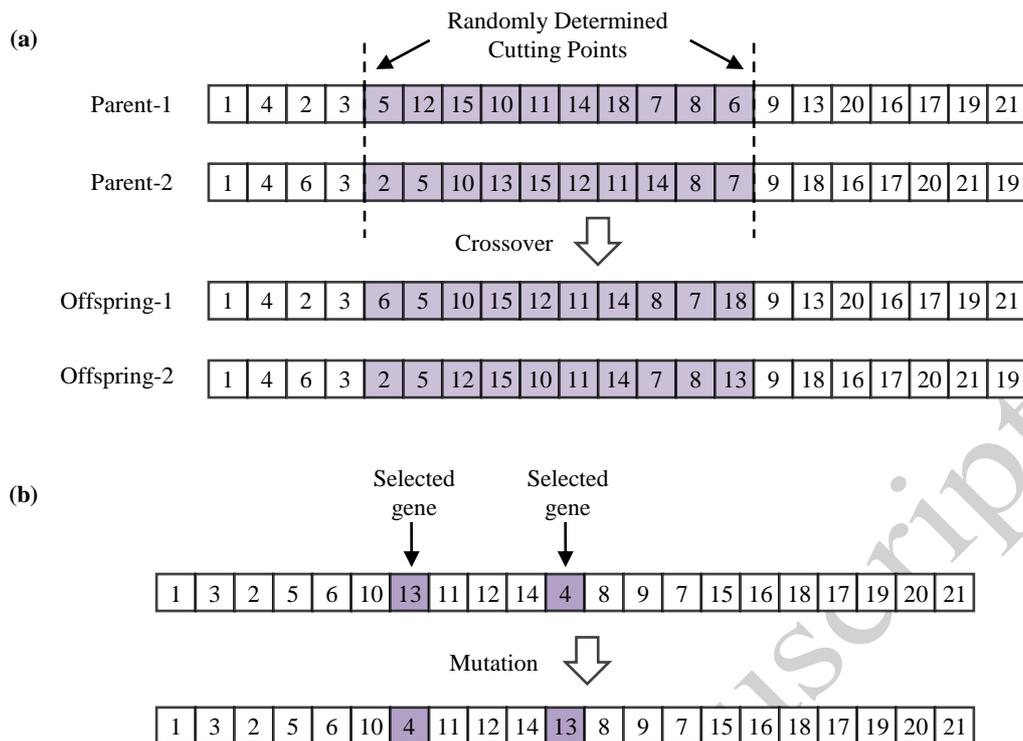


Figure 6. Illustration of (a) crossover and (b) mutation procedures.

3.5. Forming new generation

New generation is formed by comparing the fitness values of new individuals, which are obtained from crossover – mutation procedures and random generations, with existing chromosomes in the population and replacing the worst chromosomes in the population with better ones (if any).

4. Illustrative Example

To explain the running mechanism of the proposed algorithm and the encoding-decoding procedures in particular, a numerical example is given in this section. In the example, meaning of the genes, decoding procedure of the tasks, and assigning tasks to the stations can be investigated visually.

Two well-known test problems, P9 and P12 (Kim *et al.* 2000), are taken from the literature and given in Figure 7 to be considered as precedence relationships and task

processing times of two different product models (Product Model-I and Product Model-II respectively). Numbers in nodes represent task numbers while arrows between nodes symbolise precedence relationships between tasks. To make encoding-decoding procedures easier, each task number given in nodes and belonging to Product Model-II (P12) is represented by the sum of the total number of tasks belonging to Product Model-I (P9) and the original task number. For example, Task-5 of Product Model-II is represented as 14 (5+9); and Task-11 is represented as 20 (11+9). So, $n_1 + n_2 = 12 + 9 = 21$ tasks are subject to balancing. Processing time and preferred operation direction of each task, which represents the side where tasks can be assigned, are also given over each node (L, R and E denote left, right and either sides, respectively). S lists, which represent candidate tasks that can be allocated to the relevant side of each line, can be constructed as in Figure 8.

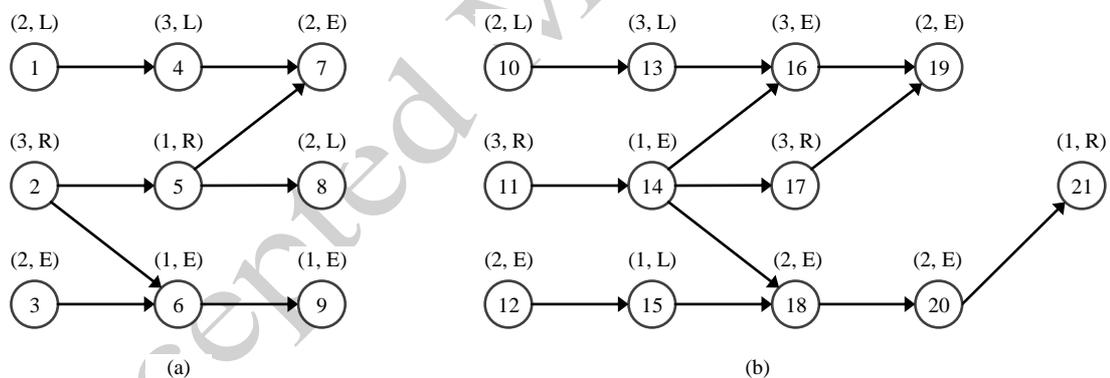


Figure 7. Precedence diagrams for the illustrative example: (a) P9, and (b) P12, adapted from Kim *et al.* (2000).

Line I	Left Side	$S1LE = \{1, 3, 4, 6, 7, 8, 9\}$
	Right Side	$S1RE = \{2, 3, 5, 6, 7, 9\}$
Line II	Left Side	$S2LE = \{10, 12, 13, 14, 15, 16, 18, 19, 20\}$
	Right Side	$S2RE = \{11, 12, 14, 16, 17, 18, 19, 20, 21\}$

Figure 8. S lists are shown on parallel two-sided assembly lines.

Task assigning process to the chromosome and workstations can be seen for the first 12 steps from Figure 9 and Figure 10, respectively. The task allocation process starts from the left side of the Line-I. If the cycle time is assumed 6 time-units for both lines, available tasks for this particular side are Task-1 and Task-3, because Task-4 – Task-9 have predecessors which have not been completed yet. Task-3 is selected randomly, and allocated to the chromosome (Step 1 in Figure 9) and to the current available station that has enough capacity (see Figure 10). Then, the station time of the current workstation is increased by the amount of assigned task's processing time. As the station time of the current workstation is larger than the station time of its mated workstation ($st(k) > st(\bar{k})$), the side is changed and available tasks are determined for the new side again. One of the available tasks (Task-2) is selected and allocated to the chromosome and to the current workstation. In Step 3, Task-6 can be initialised after completion of its predecessors, Task-2 and Task-3 (to avoid interference). When both sides of the current line are full or there is no enough capacity for assignment, the line is changed and available tasks are allocated to the adjacent line concurrently with the chromosome. This cycle continues until all tasks are assigned to the chromosome.

Step No	Line-Side	Available Tasks	Selected Task	Chromosome
1	1-L	1, 3	3	3
2	1-R	2, 6	2	3, 2
3	1-L	1, 6	6	3, 2, 6
4	1-R	5, 9	9	3, 2, 6, 9
5	1-L	1	1	3, 2, 6, 9, 1
6	1-R	5	5	3, 2, 6, 9, 1, 5
7	2-L	10, 12	10	3, 2, 6, 9, 1, 5, 10
8	2-R	11, 12	11	3, 2, 6, 9, 1, 5, 10, 11
9	2-L	12, 13, 14	13	3, 2, 6, 9, 1, 5, 10, 11, 13
10	2-R	12, 14	12	3, 2, 6, 9, 1, 5, 10, 11, 13, 12
11	2-R	14	14	3, 2, 6, 9, 1, 5, 10, 11, 13, 12, 14
12	2-L	15	15	3, 2, 6, 9, 1, 5, 10, 11, 13, 12, 14, 15

Figure 9. Task sequencing process to the chromosome.

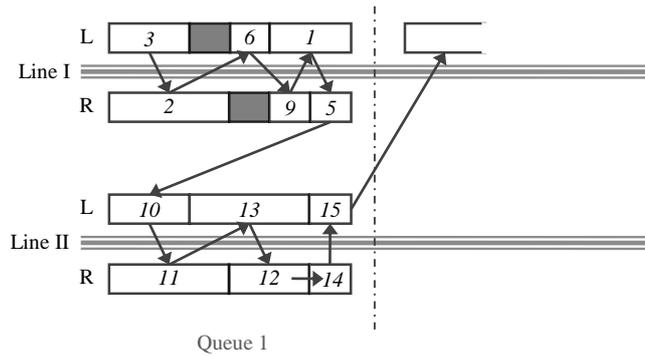


Figure 10. Task allocation process to the workstations.

Fitness evaluation of the new individuals obtained from the genetic operators and random generations are computed after the chromosome is decoded. To illustrate the decoding process, an example of decoded chromosome is demonstrated below. In the example, decoding procedure of the tasks and assigning tasks to the workstations can be investigated easily. The example chromosome given in Figure 11a can be decoded as in Figure 11b under a cycle time constraint of 6 time-units.

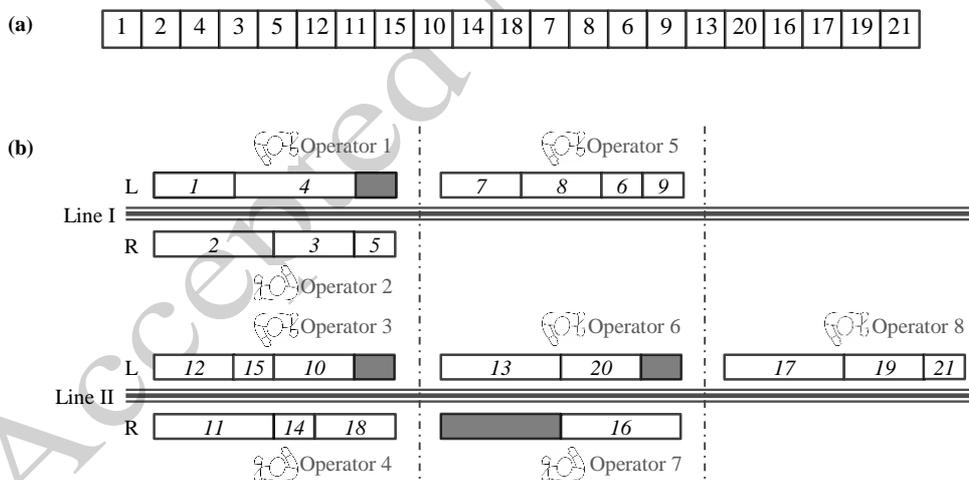


Figure 11. (a) A chromosome sample and (b) its decoded line configuration.

As can be seen in Figure 11b, eight operators are needed to assemble two different product models for the given example. On the right side of the Line-II, Task-16 can be initialised upon its predecessor task, Task-13, is completed on the left side of the line.

5. Computational Study

The proposed algorithm was coded in Java SE 7u4 environment and run on a 3.1 GHz Intel Core i5-2400 CPU 4GB RAM computer to test the performance of the proposed algorithm solving the test problems originally combined by Ozcan *et al.* (2010b). Seven original well-known problems from the literature: P9, P12 and P24 from Kim *et al.* (2000); P16, A65 and A205 from Lee *et al.* (2001); and B148 from Bartholdi (1993) (B148 was then modified by Lee *et al.* (2001)) were derived by Ozcan *et al.* (2010b) in different combinations to test the performance of tabu search algorithm in solving parallel two-sided assembly line balancing problems. In order to analyse the efficiency of the proposed approach in the current research, these test problems are solved using the proposed GA in two stages. In the first stage, the problems are solved using the cycle times provided by (Ozcan *et al.* 2010b) and the obtained results are compared with the results of Ozcan *et al.* (2010b) to have an idea about the overall performance of the proposed GA. In the second stage, test problems are solved for binary combinations of different cycle time values of Line-I and Line-II. The main objectives are (i) to observe the response of the entire system to different levels of the parallel lines' cycle times and (ii) to determine the best cycle time pair which gives the highest line efficiency.

5.1. Stage-1: Comparison with the existing results

The algorithm is run using the parameters given in Table 2 to solve the test problems given in Table 3 and the best solution is taken after three runs for each test problem. As could be seen from Table 2, used parameters may differ from one test problem to another in order to scan search space more effectively and increase the solution building capacity of the algorithm, especially in the large-sized problems, as the search space

grows exponentially with the increasing number of tasks. These parameters are chosen experimentally for a high quality solution in an acceptable period of time. Table 3 presents the test problems used by Ozcan *et al.* (2010b) along with the cycle times considered when solving these problems in the first stage of the experimental tests.

Table 2. Parameters of the proposed GA.

Test Problem	Population size	Crossover rate	Mutation rate	Number of Iterations
1 - 8	20	0.2	0.10	30
9 - 12	30	0.3	0.15	40
13 - 16	40	0.3	0.15	60
17 - 22	50	0.4	0.20	100
23 - 28	60	0.4	0.20	150
29 - 32	80	0.5	0.20	200

Table 3. Data for test problems.

Problem No	Test problems (Line I – Line II)	Number of tasks (Line I – Line II)	Cycle time (Line I – Line II)
1	P9-P9	9-9	3-3
2	P9-P9	9-9	4-5
3	P9-P12	9-12	6-6
4	P9-P12	9-12	4-7
5	P12-P12	12-12	5-5
6	P12-P12	12-12	6-7
7	P12-P16	12-16	7-16
8	P12-P16	12-16	8-21
9	P16-P16	16-16	16-16
10	P16-P16	16-16	19-21
11	P16-P24	16-24	19-35
12	P16-P24	16-24	22-40
13	P24-P24	24-24	18-18
14	P24-P24	24-24	20-24
15	P24-A65	24-65	30-490
16	P24-A65	24-65	20-544
17	A65-A65	65-65	381-381
18	A65-A65	65-65	435-435
19	A65-A65	65-65	490-544
20	A65-B148	65-148	381-408
21	A65-B148	65-148	490-459
22	A65-B148	65-148	544-510
23	B148-B148	148-148	408-408
24	B148-B148	148-148	306-357
25	B148-B148	148-148	459-510
26	B148-A205	148-205	306-1888
27	B148-A205	148-205	510-2832
28	B148-A205	148-205	255-1510
29	A205-A205	205-205	1510-1510
30	A205-A205	205-205	2832-2832
31	A205-A205	205-205	2077-2266
32	A205-A205	205-205	2454-2643

Table 4. Comparison of the obtained computational results by means of total number of required workstations.

Problem No	Independent Balancing (Line 1+Line 2)	Theoretical Minimum Number of Stations (LB)	Balancing Together	
			TS (Ozcan <i>et al.</i> 2010b)	Proposed GA
1	6 + 6	12	12	12
2	5 + 4	8	8	8
3	3 + 5	7	8	8
4	5 + 4	8	9	9
5	6 + 6	10	11	11
6	5 + 4	8	9	9
7	4 + 6	9	10	10
8	4 + 5	8	8	8
9	6 + 6	11	11	11
10	5 + 5	9	10	10
11	5 + 4	9	9	9
12	4 + 4	8	8	8
13	8 + 8	16	16	16
14	8 + 6	13	14	14
15	5 + 11	16	16	16
16	8 + 10	17	18	18
17	15 + 15	27	29	29
18	13 + 13	24	25	25
19	11 + 10	20	21	21
20	15 + 13	26	28	28
21	11 + 12	22	23	23
22	10 + 11	20	21	21
23	13 + 13	26	26	26
24	18 + 15	32	33	33
25	12 + 11	22	23	23
26	18 + 15	30	33	33
27	11 + 10	19	21	21
28	21 + 18	36	39	38
29	18 + 18	31	36	35
30	10 + 10	17	20	20
31	14 + 12	22	26	26
32	12 + 11	19	23	23

Table 4 exhibits the results (number of workstations) obtained using the proposed GA for each test problem under designated cycle time constraints. The number of stations obtained from the proposed algorithm is compared with the independent line balance of the two-sided assembly lines, the theoretical minimum number of stations (LB), and the tabu search algorithm proposed by Ozcan *et al.* (2010b) (which is the only study available in the literature and given as TS in Table 4). Obtained results are compared with respect to total number of required workstations as this is the only result reported

by Ozcan *et al.* (2010b).

Theoretical minimum number of workstations are calculated by Ozcan *et al.* (2010b). They modified simple lower bound equation proposed by Hu *et al.* (2008) for the two-sided assembly line balancing problems. As the calculation of lower bound does not take the precedence constraints (Akpınar and Bayhan 2011) into consideration, real value of the lower bound is most likely larger than the computed value, and this situation must be taken into account to measure the efficiency of the developed approach and comparison with the LB. Since the optimal number of stations cannot be less than LB, if the obtained number of stations equals to the LB, then it can be said that the obtained result is optimal (Ozcan *et al.* 2010b). As can be seen from Table 4, the proposed GA discovered optimal solutions for 9 of the 32 test problems. Moreover, GA produced one less workstation than the tabu search algorithm for the test problems 28 and 29. Therefore, it could be said that the proposed GA based approach has a promising solution capacity for the parallel two-sided assembly line balancing problems.

A Paired Two-Samples t-Test is conducted using Data Analysis tool available in Microsoft Excel™ 2010 to determine whether there is a significant difference between *independent balancing* and *together balancing* of the lines in terms of the means of number of workstations needed. The results presented in *Independent Balancing* and *Proposed GA* columns in Table 4 are subject to consideration for this statistical test. The null and alternative hypotheses are stated at the $\alpha = 0.05$ level (95%) for means of workstation numbers obtained when the lines are balanced independently (μ_I) and when the lines are balanced together using GA (μ_T) as follows:

H_0 : *There is no significant difference between the means of workstation numbers obtained by the solution strategies in favour of the alternative ($\mu_I \leq \mu_T$).*

H_1 : *Balancing lines together using the proposed GA approach significantly reduces the number of workstations needed ($\mu_I > \mu_T$).*

As seen from the hypotheses, the test is designed as one-tailed. The summary of the test results given in Table 5 (see *Test-1* column) indicates that there is significant difference in the means of workstation numbers when the lines are balanced independently ($\mu_I = 19.06, Var_I = 83.42$) and when the lines are balanced together using the proposed GA based approach ($\mu_T = 18.81, Var_T = 82.29$); $t(31) = 3.2146, p = 0.0015$. Thus, the null hypothesis is rejected with very strong evidence. These results suggest that balancing lines together allowing multi-line stations helps reduce total number of required workstations significantly.

Table 5. Results of the Paired Two-Samples t-Test for means of workstation numbers.

Paired Two-Samples t-Test	<i>Test-1</i>		<i>Test-2</i>	
	<i>Independent Balancing</i>	<i>Together Balancing (GA)</i>	<i>Together Balancing (TS)</i>	<i>Together Balancing (GA)</i>
Mean (μ)	19.06	18.81	18.88	18.81
Variance (Var)	83.42	82.29	84.63	82.29
Observations	32	32	32	32
Pearson Correlation	0.998		0.999	
Hypothesised Mean Difference	0		0	
Degrees of Freedom	31		31	
t Stat	3.2146		1.4376	
$p(T \leq t)$ one-tail	0.0015		0.0803	
t Critical one-tail	1.6955		1.6955	
$p(T \leq t)$ two-tail	0.0030		0.1606	
t Critical two-tail	2.0395		2.0395	

Using the results proposed in the relevant columns of Table 4, another Paired Two-Samples t-Test is performed to determine whether there is a significant difference between the means of workstation numbers found by TS (Ozcan *et al.*) and proposed GA. The null and alternative hypotheses stated at the $\alpha = 0.05$ level (95%) for means of workstation numbers obtained using TS (μ_{TS}) and GA (μ_{GA}) are as follows:

H_0 : *There is no significant difference between the means of workstation numbers obtained by the solution strategies in favour of the alternative ($\mu_{TS} \leq \mu_{GA}$).*

H_1 : GA algorithm finds better solutions than TS when balancing parallel two-sided assembly lines ($\mu_{TS} > \mu_{GA}$).

Based on the summary of the test results presented in Table 5 (see *Test-2* column), there is no strong evidence to reject the null hypothesis at $\alpha = 0.05$. Therefore, it is not possible to argue that GA ($\mu_{GA} = 18.81, Var_{GA} = 82.29$) performs significantly better than TS ($\mu_{GA} = 18.88, Var_{GA} = 84.63$) at this confidence level; $t(31) = 1.4376, p = 0.08$. However, it could be argued that GA finds significantly better solutions than TS if the test was performed at $\alpha = 0.1$, and it can be clearly seen from Table 4 that GA finds quite promising results.

5.2. Stage-2: Solutions for various cycle time situations

Now, we can proceed to the second stage of the computational tests assuming that the performance of the proposed GA based algorithm is sufficient enough. In this stage, the test problems given above are solved using the proposed GA (with the same GA parameters used in the previous subsection) by considering different cycle times for the lines. Four levels are determined for cycle time of each line in each test problem and the problems are solved under the constraints of these cycle time combinations. Considered cycle times for Line-I and Line-II, calculated common cycle time (C), and obtained number of stations (K) are given in Table 6 for different test cases. The *LE* column reports the computed system efficiency based on the obtained number of workstations. This value is obtained via dividing total needed time to perform all tasks on the lines by the total available time of the utilised system (see Equation 9, the definitions of the used symbols have already been given in Section 2.2).

$$LE = \frac{\sum_{h=1}^H \frac{C}{C_h} \sum_{i=1}^{n_h} t_{hi}}{K \times C} . \quad (9)$$

Table 6. Computational results for different cycle time levels of the lines.

P9-P9													
T. Task Times		Cycle Time of Line-II											
P9	P9	4			5			6			7		
17	17	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	3	12	11	0.90	15	10	0.91	6	9	0.94	21	9	0.90
	4	4	9	0.94	20	9	0.85	12	8	0.89	28	8	0.83
	5	20	8	0.96	5	8	0.85	30	7	0.89	35	7	0.83
	6	12	8	0.89	30	7	0.89	6	6	0.94	42	6	0.88

P9-P12													
T. Task Times		Cycle Time of Line-II											
P9	P12	6			8			10			12		
17	25	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	4	12	10	0.84	8	8	0.92	20	8	0.84	12	7	0.90
	6	6	8	0.88	24	7	0.85	30	6	0.89	12	6	0.82
	8	24	8	0.79	8	6	0.88	40	6	0.77	24	6	0.70
	10	30	7	0.84	40	6	0.80	10	5	0.84	60	5	0.76

P12-P12													
T. Task Times		Cycle Time of Line-II											
P12	P12	6			8			10			12		
25	25	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	5	30	11	0.83	40	10	0.81	10	9	0.83	60	8	0.89
	7	42	9	0.86	56	8	0.84	70	7	0.87	84	7	0.81
	9	18	8	0.87	72	7	0.84	90	7	0.75	36	6	0.81
	11	66	8	0.80	88	7	0.77	110	6	0.80	132	6	0.73

P12-P16													
T. Task Times		Cycle Time of Line-II											
P12	P16	16			18			20			22		
25	82	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	8	16	10	0.83	72	9	0.85	40	9	0.80	88	8	0.86
	10	80	9	0.85	90	9	0.78	20	8	0.83	110	7	0.89
	12	48	9	0.80	36	8	0.83	60	8	0.77	132	7	0.83
	14	112	8	0.86	126	8	0.79	140	8	0.74	154	6	0.92

P16-P16													
T. Task Times		Cycle Time of Line-II											
P16	P16	17			19			21			23		
82	82	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	14	238	13	0.82	266	13	0.78	42	13	0.75	322	12	0.79
	16	272	12	0.83	304	12	0.79	336	12	0.75	368	10	0.87
	18	306	12	0.78	342	12	0.74	126	11	0.77	414	11	0.74
	20	340	11	0.81	380	11	0.77	420	11	0.73	460	10	0.77

P16-P24													
T. Task Times		Cycle Time of Line-II											
P16	P24	25			27			29			31		
82	140	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	20	100	11	0.88	540	11	0.84	580	11	0.81	620	11	0.78
	22	550	11	0.85	594	10	0.89	638	10	0.86	682	10	0.82
	24	600	10	0.90	216	10	0.86	696	10	0.82	744	10	0.79
	26	650	10	0.88	702	10	0.83	754	10	0.80	806	10	0.77

Table 6 (continued).

P24-P24													
T. Task Times		Cycle Time of Line-II											
P24	P24	19			21			23			25		
140	140	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	18	342	17	0.89	126	16	0.90	414	15	0.92	450	15	0.89
	20	380	16	0.90	420	16	0.85	460	15	0.87	100	15	0.84
	22	418	15	0.92	462	15	0.87	506	15	0.83	550	14	0.85
	24	456	15	0.88	168	14	0.89	552	14	0.85	600	13	0.88

A65-A65													
T. Task Times		Cycle Time of Line-II											
A65	A65	385			425			465			505		
5099	5099	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	360	27720	32	0.86	30600	31	0.84	11160	29	0.87	36360	28	0.87
	390	30030	30	0.88	33150	28	0.90	12090	28	0.86	39390	27	0.86
	420	4620	29	0.88	35700	28	0.86	13020	27	0.86	42420	26	0.86
	450	34650	29	0.85	7650	26	0.90	13950	26	0.86	45450	25	0.86

A65-B148													
T. Task Times		Cycle Time of Line-II											
A65	B148	375			400			425			450		
5099	5024	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	360	9000	32	0.86	3600	31	0.86	30600	30	0.87	1800	30	0.84
	380	28500	32	0.84	7600	30	0.87	32300	29	0.87	17100	29	0.85
	400	6000	31	0.84	400	30	0.84	6800	29	0.85	3600	28	0.85
	420	10500	30	0.85	8400	29	0.85	35700	28	0.86	6300	28	0.83

B148-B148													
T. Task Times		Cycle Time of Line-II											
B148	B148	300			350			400			450		
5024	5024	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	325	3900	38	0.85	4550	34	0.88	5200	33	0.85	5850	32	0.83
	375	1500	34	0.89	5250	32	0.87	6000	30	0.87	2250	30	0.82
	425	5100	34	0.84	5950	31	0.84	6800	29	0.84	7650	28	0.82
	475	5700	33	0.83	6650	29	0.86	7600	28	0.83	8550	26	0.84

A205-A205													
T. Task Times		Cycle Time of Line-II											
A205	A205	1550			1850			2150			2450		
23345	23345	C	K	LE	C	K	LE	C	K	LE	C	K	LE
Cycle Time of Line-I	1475	91450	40	0.77	109150	36	0.79	126850	34	0.78	144550	34	0.75
	1850	57350	36	0.77	1850	32	0.79	79550	32	0.73	90650	30	0.74
	2225	137950	34	0.75	164650	31	0.75	191350	28	0.76	218050	27	0.74
	2600	80600	32	0.75	96200	29	0.74	111800	27	0.73	127400	25	0.74

'T.Task Times' column gives the sum of all task times for the relevant problem.

Line efficiency is a well-known term which is commonly used as a measure of the obtained solution's quality regardless of the tackled line configuration and problem type. Therefore, the proximity of a line system's efficiency to '1' could be considered as an indicator whether this system is well balanced or not. If the efficiency equals to '1',

this means that there is no idle time on the line. However, this is hardly possible in such systems due to unsmooth task times and cycle time differences between the lines.

Although the optimality of the solutions cannot be guaranteed, our findings indicate that near-optimal solutions can be obtained very quickly for even large-sized instances. As could be seen from the obtained results, different solutions are obtained with different line efficiency values corresponding to the binary combinations of the cycle time levels of the parallel lines for the same test problem. The proposed algorithm finds high quality solutions over 85% efficiency for the entire test problems studied, except the case A205-A205. The best obtained efficiency result (96%) belongs to the problem P9-P9 and is obtained with the cycle time combination of 5 and 4 for Line-I and Line-II, respectively. The next best result, 92%, is obtained for the problems P9-P12, P12-P16, and P24-P24. As a result of the NP-Hard characteristic of the studied problem, it is reasonable that the efficiency value obtained for the largest problem (79% for A205-A205) is lower than those for the small-sized ones as expected. However, even this result could be quite reasonable for real world scenarios.

Obtained efficiency values across two dimensions (the cycle time of Line-I and the cycle time of Line-II) are also plotted as a surface graph and are depicted in the Appendices. Thus, readers can see the best provided cycle time combination, which gives the highest efficiency, for each case easily. Also, managers can pick up the *number of workstation – cycle time* combination that fits their organisations and model demands from the provided results.

6. Conclusions

The simple assembly line balancing problem is an NP-hard class of combinatorial problem, as shown by Wee and Magazine (1982). Since the PTALBP is a much more complex version of the simple assembly line balancing problem, it is also NP-Hard. The

solution space grows exponentially as the number of tasks increases, which means that obtaining an optimal solution when the problem size increases is very difficult (Kalayci and Gupta 2014). It is the major reason why a considerable amount of researches in the literature strives to develop heuristics and meta-heuristics instead of exact methods to solve the assembly line balancing problems.

We developed the first mathematical model for a recently introduced production planning problem, *parallel two-sided assembly line balancing problem*, and proposed an alternative possible approach for the solution of the problem. Since the problem is very complex and the size of the problems that can be solved in an acceptable amount of time is drastically limited, we applied GA to the PTALBP, which is the first GA based approach to solve such a problem, and have obtained very encouraging results. To assess the performance of the algorithm, a set of test problems, previously combined and solved by Ozcan *et al.* (2010b), are solved and obtained results are compared with Ozcan *et al.* (2010b). Although the complexity of the problem is higher than other configurations of assembly lines (*i.e.* one-sided straight assembly lines), computational results demonstrate that the performance of the proposed algorithm is sufficient. Moreover, the effect of different cycle time situations on the efficiency of the overall line system is also studied for a parallel line system for the first time in the literature. For this aim, parallel lines are balanced for different combinations of their different cycle time levels and the efficiency of the entire line system is reported along with the total number of required stations for each case. Obtained line efficiencies are plotted as surface charts across cycle times of the parallel lines to make analyses easier. Thus, line managers can easily pick up the *number of workstations - cycle time* pair that suits their company best.

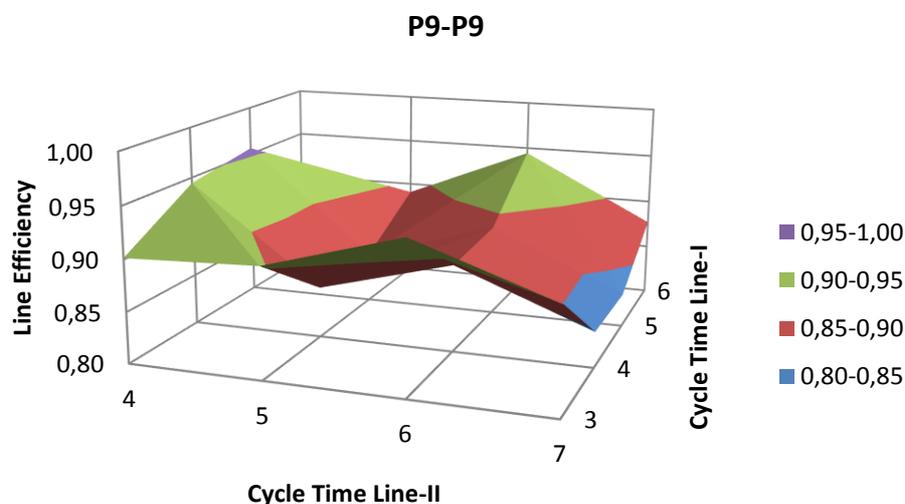
This study makes it clear that more research is needed to fill in the gap in the literature on minimising cycle time of the parallel two-sided lines as well as total number of required workstations. Hybrid meta-heuristics and/or hyper-heuristics might also be proposed to increase the solution capacity of the algorithm; or exact solution procedures may be developed to solve the PTALBP, even not the large-sized instances. In addition, workload smoothness between workstations and lines may be of interest for future studies with some more realistic conditions of real applications (*i.e.* zoning constraints, task synchronisation constraints, positional constraints, *etc.*).

Acknowledgement

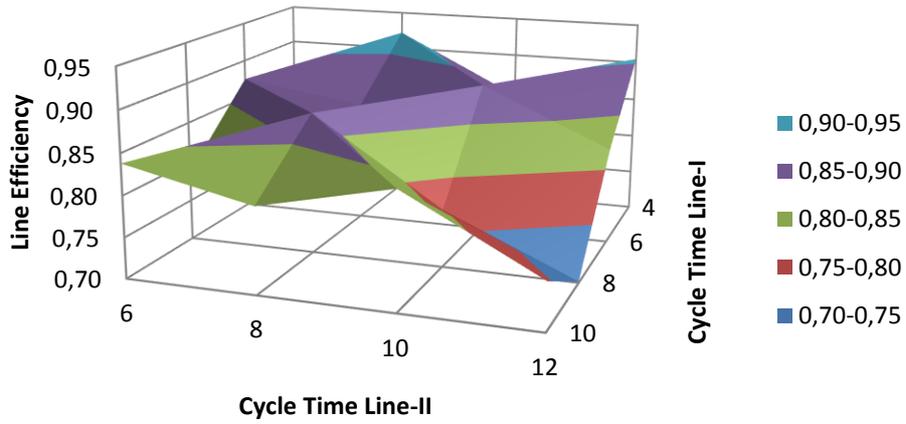
The first author gratefully acknowledges the financial support from the *Balikesir University* and the *Turkish Council of Higher Education* during his Ph.D. at the *University of Exeter* in England. Both authors are grateful to anonymous reviewers for their valuable comments and suggestions to improve the quality of the paper.

Appendices

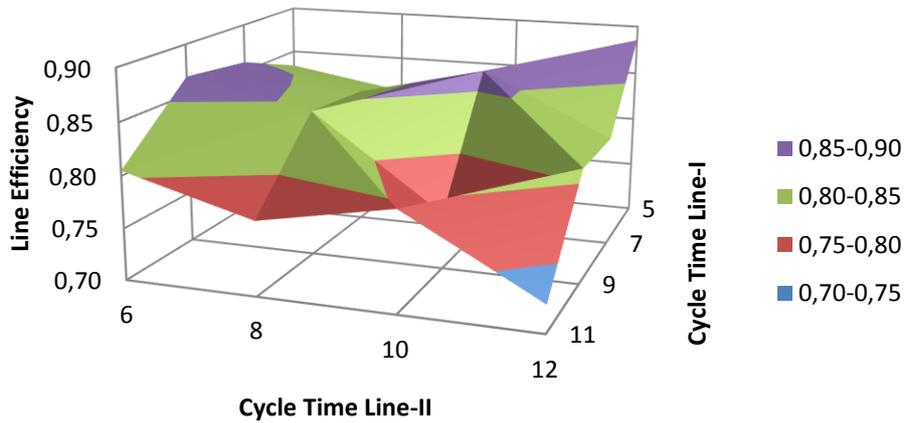
A.1. Surface charts of the line efficiency values across the cycle times of Line-I and Line-II for the test problems.



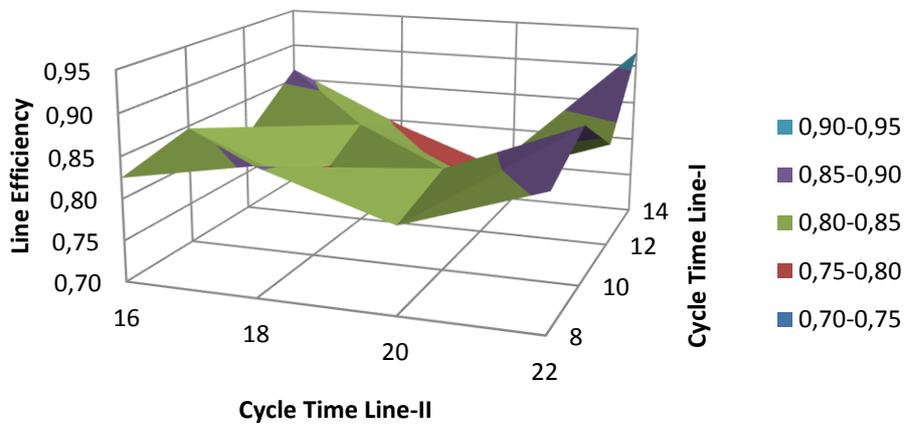
P9-P12



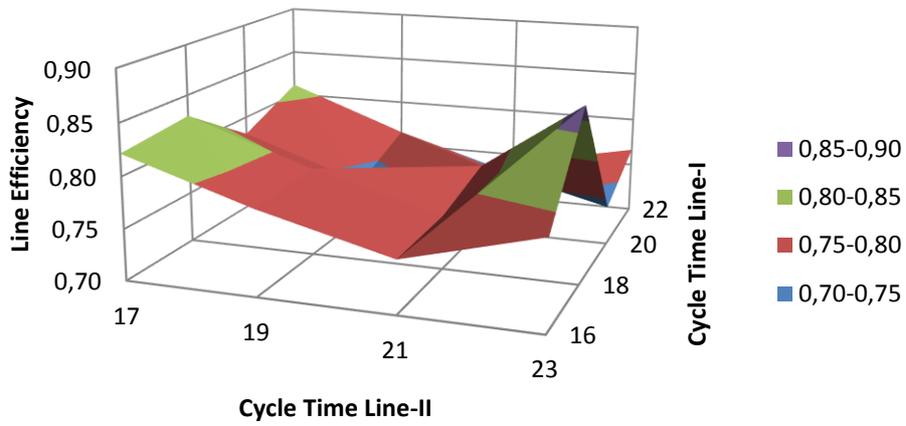
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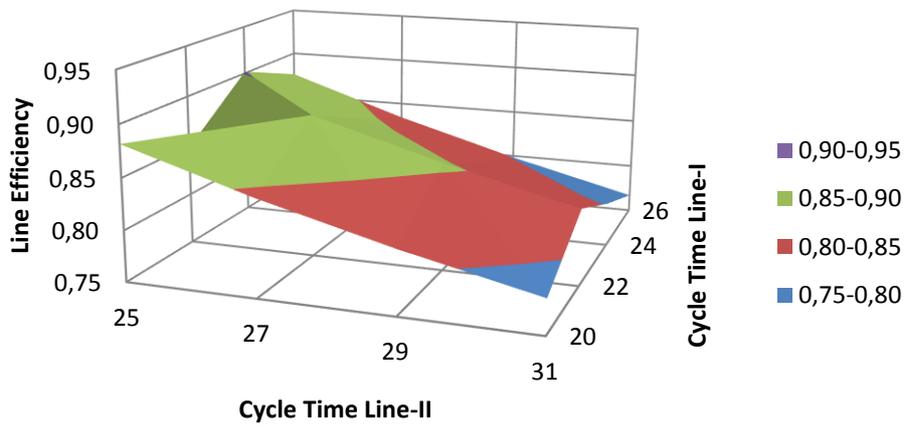
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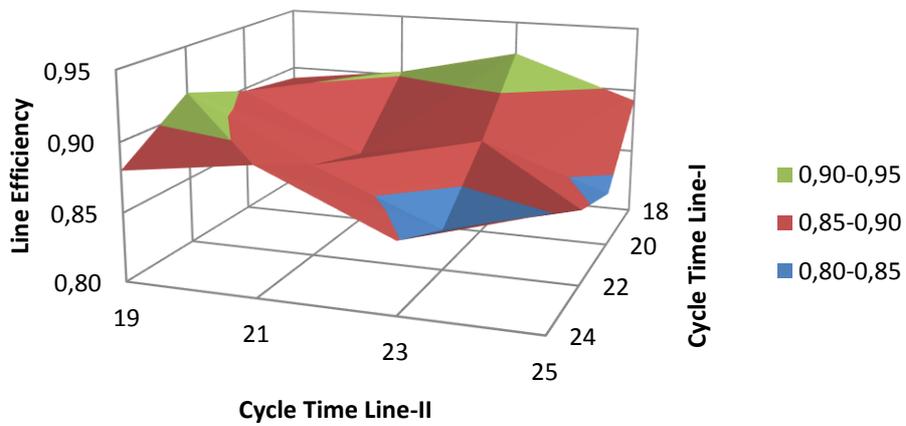
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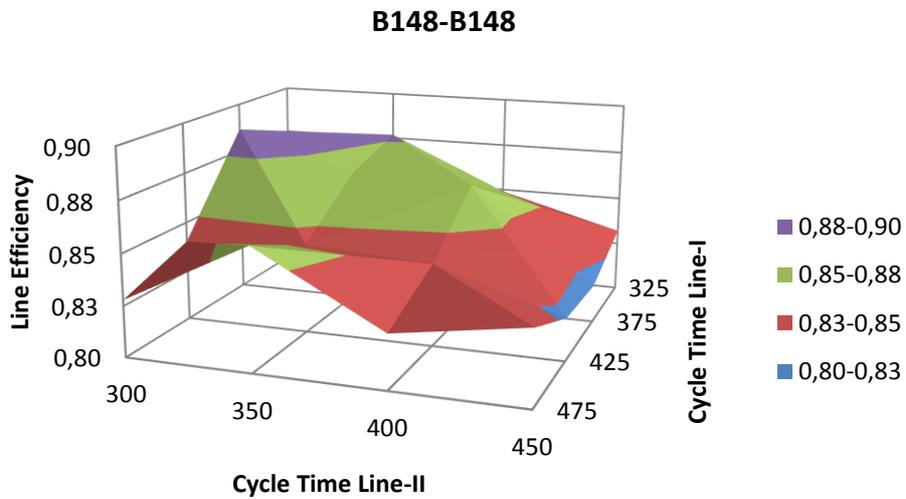
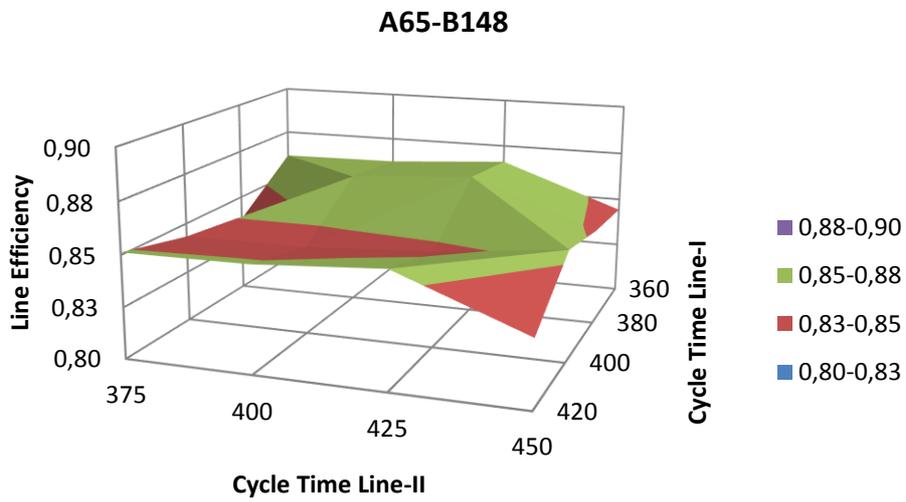
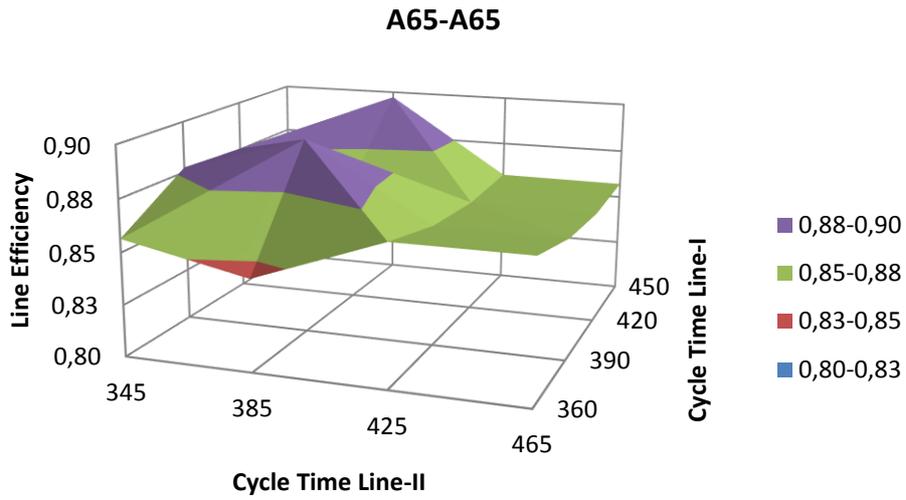


P16-P24

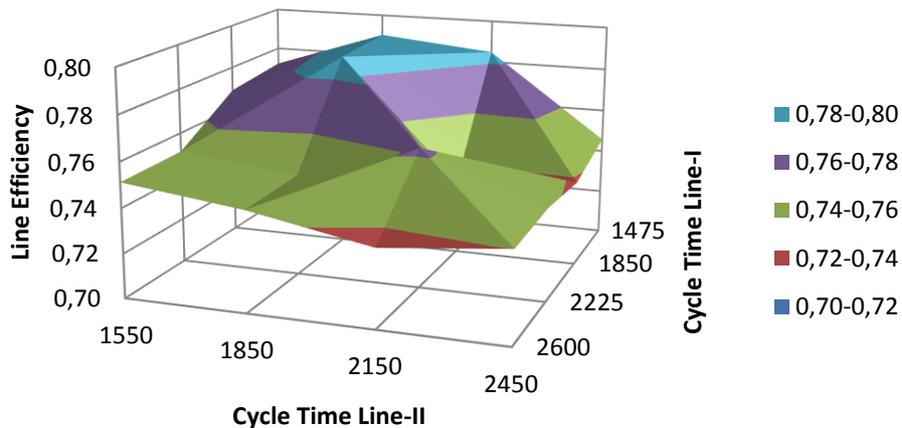


P24-P24





A205-A205



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