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On Optimizing a Demand Responsive Transport with an Evolutionary Multi-Objective Approach

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Abstract—This paper deals with a dial-a-ride problem with time windows applied to a demand responsive transport service. An evolutionary approach as well as new original representation and variation operators are proposed and detailed. Such mechanisms are used with three state-of-the-art multi-objective evolutionary algorithms: NSGA-II, IBEA and SPEA2. After introducing the general problem, the solution encoding and the algorithm mechanisms are depicted. The approach is assessed by applying the algorithms to both random and realistic dial-aride instances. Then a statistical comparison is provided in order to highlight the most suited evolutionary algorithms to optimize real-life transportation problems.

Index Terms—Dial-a-ride, Demand responsive transport, Multi-objective, Evolutionary algorithm, Optimization

I. INTRODUCTION

The demand responsive transport (DRT) is a transport service which involves the satisfaction of a set of travel requests. A request corresponds to a customer who wants to be carried from a pick-up location to a delivery at a predetermined time. In its usual form, DRT is related to the dial-a-ride problem (DARP) or to the vehicle routing problem (VRP), which both consist in optimizing the vehicles routes by reducing the number of vehicles and the travel durations [1], [2]. Nevertheless, the main differences between the DARP and the VRP consist of the precedence constraints imposed by the customer journeys and in their tolerance to the delays (quality of service [3]). A DRT service is a specific case of the DARP, the latter being the academical formulation of routing with passengers.

Given that the DARP is *NP*-hard [1], we propose to use an Evolutionary Algorithm (EA), which is known to efficiently cope with scalable problems. Moreover, EAs can produce multiple distinct solutions in a single run which could help the decision making. Besides, the DARP involves antagonist objectives which have to be optimized concurrently. Since using an EA paired to a Pareto approach has already been shown to efficiently solve multi-objective real-life problems [4], [5], a Pareto approach is preferred in our application in order to not hierarchize the criteria under study.

Although a lot of metaheuristic methods exist for solving the DARP [6], [1], EA are generally used as single-criterion optimization methods aggregating multiple objectives in order to make the computation easier [7], [8], [9]. As far as we know, only Chevrier [10] used a Pareto-based three-objectives model to cope with the DARP with time window (DARPTW).

In this paper, a new encoding mechanism based on a twodimensional representation as well as original operators are proposed for the DARPTW. They are integrated in three state-of-the-art algorithms: the Non dominated Sorting Genetic Algorithm II (NSGA-II) [11], Strength Pareto EA 2 (SPEA-2) [12] and Indicator Based EA (IBEA) [13]. Then they are applied to benchmark instances and pairwise compared.

The paper is organized as follows. The problem formulation is described in Section II. In Section III, the evolutionary multi-objective principles are introduced. Then we present the proposed encoding used to represent the DARP, as well as the variation operators, in Section IV. Experimental results for Chevrier's instances [10] are presented and discussed in Section V. Finally, Section VI concludes the paper.

II. PROBLEM OVERVIEW

A. Definitions

First, we define a set of basic notations useful for the remainder of the paper. Let r be a request and R the set of the requests: $r \in R$. Let V denote the set of pick-up and delivery locations. A request r is defined by: r^+, r^- the (resp.) pick-up and delivery locations: $\{r^+, r^-\} \in V; h_{r^+}$ the desired pick-up time; w_r the number of customers to be carried from r^+ to r^- . The travel durations are stored in a matrix M. Let x, y be two points and $t_{x \to y}$ the travel duration from x to $y: t_{x \to y} = M_{xy}$. Note that the matrix M is not necessarily symmetrical: $M_{xy} \neq M_{yx}$. Given a travel duration, we can deduce a theoretical delivery time $h_{r^-} = h_{r^+} + t_{r^+ \to r^-}$.

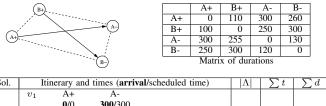
B. Objectives

In the DRT service under study, we aim at minimizing three objectives. The first (φ_1) is *economic* and consists in minimizing the number of required vehicles. The second corresponds to the duration reduction of the vehicle journeys (φ_2) which could correspond to an *environmental* objective in so far as we look to limiting the emission of pollutants, and also in expectation of a hypothetical 'carbon tax'. The last objective minimizes the likely delays (*quality of service*: φ_3). In the remainder of the paper, Λ denotes the set of required vehicles, t a duration, and d a delay.

C. Example

In the example of Figure 1, a DRT service must transport two customers. Customer A wants to go from Point A+ to Point A- at $h_{A+}=0$, while Customer B wants to be carried from B+ to B- at $h_{B+} = 100$. The matrix of durations indicates the 'point to point' durations (a line index indicates the origin and a column the destination). Note that the service proposed here is not 'door-to-door' because it is more usual to pick-up and deliver customers at predefined stops (those of the public transport company)

Several solutions could be designed according to the objectives under consideration. Each detailed solution has advantages and drawbacks. Hence, Solution S_1 requires two vehicles $(|\Lambda| = 2)$, whereas Solutions S_2 and S_3 need only one vehicle. According to the economic point of view, S_2 and S_3 are better than S_1 . However, this latter has no delay ($\sum d = 0$) contrary to S_2 and S_3 which have a few delay (resp. 140 and 150). If we consider the quality of service only, S_1 would be better than S2 and S3. In the same way, the total durations spent in traveling $(\sum t)$ indicate that S_2 would be the best solution in this case. Thus, these three solutions are incomparable if we want to optimize all objectives concurrently [4]. Given that the research of an optimal solution in a multi-criteria problem is intractable [5], we propose a multi-objective algorithm based on a Pareto approach.



S_1	v_2	0/0 B+ 100/100	300 /300 B- 400 /400			2	600	0
S_2	v_1	A+ 0/0	B+ 110/100	B- 410 /400	A- 530/400	1	530	140
S_3	v_1	A+ 0/0	B+ 110/100	A- 360 /300	B- 490 /400	1	490	150

Fig. 1. Example of a DARP instance with two customers and three solutions $S_1, S_2, S_3.$

D. Introduction of a delay tolerance

A usual DRT service uses the tolerance of the customers to accept more or less delays. Making detours allows a vehicle to group the customers more easily while producing a few delays. In this perspective, we introduce a coefficient of relaxation k_r applied to the travel duration to define a maximal delivery time. Let $t'_{r^+ \to r^-}$ be the slackened travel duration: $t'_{r^+ \to r^-} = k_r \cdot t_{r^+ \to r^-}$. Consequently, the maximal delivery time h'_{r^-} is defined as follows: $h'_{r^-} = h_{r^+} + t'_{r^+ \rightarrow r^-}$.

E. Introduction of time windows

In the case of VRP, time windows are generally used as time slots to make delivery easier. In the case of DARP, a time window facilitates the picking-up when a travel duration slightly exceeds the picking-up time. With such a mechanism, it is easier to bring together customers into the same vehicles. So, a time window in a point r^+ is denoted tw_{r^+} and is proportional to the theoretical travel duration to the point r^{-} : $tw_{r^+} = k_w t_{r^+ \rightarrow r^-}$, where k_w is a coefficient indicating the percentage of the duration allocated to the time window.

F. Constraints

The flexibility is introduced by using relaxation and time windows while authorizing delays on the travels. Nevertheless, for limiting delays and for keeping a good quality of service, the likely delays must be limited by adding a set of constraints. Firstly, it is necessary to define what a feasible travel is. We denote x, y two arbitrary points to be potentially connected and H_x the effective starting time at Point x. Indeed, there may be some accumulated delays that H_x takes into account. Two cases may happen:

- if y is a picking-up point then the travel from x to y is feasible iff $H_x + t_{x \to y} \leq h_y + tw_y$
- if y is a delivery point then the travel from x to y is feasible iff $H_x + t_{x \to y} \le h'_y$

Note that H is evaluated after each travel between two points and depends on the travel durations and whether the vehicle arrives before or after a theoretical starting time. Therefore, two cases may arise for determining H_y , which corresponds to the effective time in Point y coming from Point x:

- $H_x + t_{x \to y} < h_y \Rightarrow H_y = h_y;$ $H_x + t_{x \to y} \ge h_y \Rightarrow H_y = H_x + t_{x \to y}$

After calculating the effective visiting time, it is possible to evaluate the delay at a delivery point. A delay d corresponds to the difference between H and the theoretical arrival time, thus in a delivery point y:

$$d_y = \begin{cases} H_y - h_y & \text{if } H_y > h_y \\ 0 & \text{otherwise.} \end{cases}$$

III. EVOLUTIONARY MULTI-OBJECTIVE OPTIMIZATION

A. Multi-objective Optimization

A general Multi-objective Optimization Problem (MOP) can be defined by a set of n objective functions (f_1, f_2, \ldots, f_n) , a set X of feasible solutions in the decision space, and a set Z of feasible points in the objective space. Without loss of generality, we here assume that each objective function is to be minimized. To each solution $x \in X$ is assigned an objective vector $z \in Z$ on the basis of the vector function $f: X \to Z$ with $z = f(x) = (f_1(x), f_2(x), \dots, f_n(x))$ as illustrated by Figure 2. An objective vector $z \in Z$ is said to dominate¹ another objective vector $z' \in Z$ iff $\forall i \in \{1, 2, ..., n\}, z_i \leq z'_i$ and $\exists j \in \{1, 2, ..., n\}$ such as $z_j < z'_j$. An objective vector $z \in Z$ is said to be *non-dominated iff* there does not exist another objective vector $z' \in Z$ such that z' dominates z. A solution $x \in X$ is said to be *efficient* if its mapping in the objective space results in a non-dominated point. The set of all efficient solutions is the *efficient set*, denoted by X_E . The

¹We will also say that a decision vector $x \in X$ dominates a decision vector $x' \in X$ if f(x) dominates f(x').

set of all non-dominated vectors is the *Pareto front*, denoted by Z_N . A possible approach in MOP solving is to find the minimal set of efficient solutions, *i.e.* one solution $x \in X_E$ for each non-dominated vector $z \in Z_N$ such as f(x) = z. However, generating the entire efficient set is usually infeasible due to the complexity of the underlying problem. Therefore, the overall goal is often to identify a good approximation of it. EAs are commonly used to this end as they are able to find multiple and well-spread non-dominated solutions in a single simulation run [4].

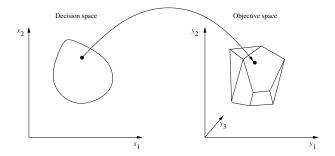


Fig. 2. Representation of a solution (x_1, x_2) in the decision space and the corresponding values in the objective space: $(y_1, y_2, y_3) = f(x_1, x_2)$.

B. Evolutionary Multi-objective Algorithms

1) NSGA-II: [11] is probably the most widely used multiobjective resolution method. At each NSGA-II generation, solutions from the current population are ranked into several classes. Individuals mapping to vectors from the first front all belong to the best efficient set; individuals mapping to vectors from the second front all belong to the second best efficient set; and so on. Two values are then assigned to the population members. The first one corresponds to the rank the corresponding solution belongs to, and represents the quality of the solution in terms of convergence. The second one, the *crowding distance*, consists in estimating the density of solutions surrounding a particular point of the objective space, and represents the quality of the solution in terms of diversity. A solution is said to be better than another one if it has a best rank value, or in case of equality, if it has the best crowding distance. The selection strategy is a deterministic tournament between two random solutions. At the replacement step, only the best individuals survive, with respect to a predefined population size. Furthermore, note that, in addition to the original NSGA-II, we add an external population, the so-called archive, in order to store the whole set of potentially efficient solutions found during the search.

2) *IBEA*: [13] is a more recent method that is a good illustration of the new trend dealing with indicator-based search, and started to become popular in recent years. The main idea of IBEA is to introduce a total order between solutions by means of a binary quality indicator. Its fitness assignment scheme is based on a pairwise comparison of solutions from the current population with regards to an arbitrary indicator *I*. To each individual *x* is assigned a fitness value F(x) measuring the 'loss in quality' if *x* was removed from the current

population P, *i.e.* $F(x) = \sum_{x' \in P \setminus \{x\}} (-e^{-I(x',x)/\kappa})$, where $\kappa > 0$ is a user-defined scaling factor. Different indicators can be used for such a purpose, and we choose to use the binary additive ϵ -indicator $(I_{\epsilon+})$ as defined in [13]. $I_{\epsilon+}(x,x')$ gives the minimum value by which a solution $x \in X$ has to or can be translated in the objective space to weakly dominate another solution $x' \in X$. Selection for reproduction consists of a binary tournament between randomly chosen individuals. Selection for replacement consists in iteratively removing the worst solution from the current population until the required population size is reached; fitness information of the remaining individuals is updated each time there is a deletion. Similarly, we added an archive to store non-dominated solutions in order to prevent their loss during the stochastic search process.

3) SPEA2: [12] is an extension of SPEA, where an improved fitness assignment strategy is proposed. It intrinsically handles an internal archive of fixed size that is used during the selection step to create offspring solutions. At a given iteration of SPEA2, to each population and archive member x is assigned a strength value S(x) representing the number of solutions it dominates. Then, the fitness value F(x) of solution x is calculated by summing the strength values of all individuals solution x currently dominates. Additionally, a diversity preservation strategy, based on a nearest neighbor technique, is incorporated. The selection step consists of a binary tournament with replacement applied on the internal archive only. At last, given that the SPEA2 archive has a fixed size storage capacity, a bounding mechanism, based on fitness and diversity information, is used when the size of the non-dominated set is too high. On the contrary, when the size of the non-dominated set is too small, some dominated solutions are allowed to be incorporated. As well, we added an external archive to store the whole set of non-dominated solutions found during the search.

IV. APPLICATION TO THE DARPTW

A. Solution encoding and initialization

1) Representation: The main idea of the representation is to simplify the reading of the vehicle routes. So, we use a two-dimensional representation (not a matrix but a vector of vectors), in which a gene corresponds to a single vehicle route (one vector). The sequence of data literally indicates the order of the points traversed by a vehicle. In order to ensure the points precedence, a cell indicates a request identifier and not the point itself in such a way that we can retrieve the associated point by counting the number of times the corresponding request identifier is encountered. If a request r appears for the first time, it is necessarily the pick-up point r^+ , otherwise it is the delivery point r^- . Given that there are two points per request, a gene has necessarily an even number of cells.

The examples of Figures 3(b,c,d) are solutions to a DARPTW instance (Fig. 3(a)). For each solution, the associated encoding is depicted. Each gene *i* is a vector translating the route of vehicle v_i . The order of the cell values indicates the sequence of the points visited by a vehicle, such as in the

example 3(b), where the sequence (A, B, A, C, B, C) is the counterpart of the path $(A^+, B^+, A^-, C^+, B^-, C^-)$.

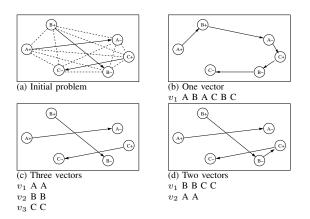


Fig. 3. Examples of solution encoding: (a) Initial problem with three customers A, B, C, (b) a solution with one vehicle (one vector), (c) a solution with three vehicles (three vectors), (d) a solution with two vehicles (two vectors).

2) Population initialization: The population is composed of N feasible solutions. To be sure of their feasibility we assign one request per vehicle, so that $|\Lambda| = |R|$. Hence, the previous DARP instance (Fig. 3(a)) requires three vehicles and the population $P = \{S_i\}$ could be initialized as follows:

	S_1	S_2	S_3	S_4				
v_1	A A	CC	A A	BB				
v_2	CC	A A	BB	A A				
v_3	BВ	ВВ	СС	CC				

B. Evaluation

The evaluation function $\Phi = (\varphi_1, \varphi_2, \varphi_3)$ applied to a solution S computes three values corresponding to the respective objectives:

- $\varphi_1 = \min |\Lambda|;$
- $\varphi_2 = \min \sum_{v \in \Lambda} t_v$, that is the minimization of the sum of each vehicle route duration t_v (t_v corresponds to the amount of each travel duration between each point visited by Vehicle v);
- $\varphi_3 = \min \sum_{v \in \Lambda} d_v$, that is the minimization of the amount of each vehicle delay d_v (d_v equals to the sum of each delay at arrivals (delivery points)).

If a solution is not feasible and marked as wrong, its score is penalized in such a way that the solution is unlikely to be kept in the next population.

C. Crossover

The crossover aims at producing new individuals (offspring) from two individuals of the current population (parents). Let P_1, P_2 be two parents in the population and C_1, C_2 two solutions built from P_1, P_2 . C_1 (resp. C_2) is the copy of P_1 (resp. P_2) with elements from P_2 (resp. P_1).

After creating C_1 , the next step consists in randomly choosing a gene of P_2 at position λ (vector v_{λ}). v_{λ} contains the values that will be reassigned to C_1 . But, in order to avoid duplicated data, the values of C_1 matching those of v_{λ} are removed. Then, the data of v_{λ} can be inserted in Gene λ at a random position α of C_1 .

In the example of Figure 4, Vector v_1 of P_2 is randomly chosen ($\alpha = 1$) and the values (C C) have to be assigned to the counterpart vector of C_1 . These values are randomly inserted into Vector v_1 of C_1 and removed from Vector v_2 . If the solutions to be crossed do not have the same size, *i.e.* the vector to be filled does not exist, a new vector is created before filling it. Producing C_2 follows the same process by inverting P_1 and P_2 .

$$\begin{array}{c} \begin{array}{c} \begin{array}{c} \text{Copying } P_1 \text{ and choosing } v_1 \\ \hline C_1 = P_1 \\ P_2 \\ P_2 \\ P_3 \\ P_3 \end{array} \xrightarrow{P_2} \begin{array}{c} C \ C \ D \ D \\ B \ B \end{array} \xrightarrow{P_2} \begin{array}{c} \begin{array}{c} \text{Assigning } (C \ C) \text{ to } v_1 \\ \hline C_1 \\ \hline C_1 \\ \hline B \ A \ C \ C \\ D \ D \end{array} \end{array}$$

Fig. 4. Example of crossover of two solutions P_1, P_2 for producing a new individual C_1 .

D. Mutation

1

The mutation generally aims at bringing diversity into a population and prevents the population to converge towards local optima. Our operator mixes two mechanisms, each one used in half cases:

- an analogy to the well-known **2-OPT** exchange operator, which reverses a sequence of values (random size) of a randomly chosen gene (Fig. 5(a)). For example a sequence (C A C B) becomes (B C A C);
- a shifting of a random customer from a vector to another (Fig. 5(b)), the pair values are randomly inserted in the host vector.

(a) Analogy to the 2	2-OPT opera	tor
C_1		C'_1
BACACB	\Rightarrow	BABCAC
DD		D D
88		
(b) Shifting of one of	customer	
$\overline{C_1}$	customer	
<u> </u>	customer ⇒	$\frac{C_1''}{A C A C}$

Fig. 5. Example of mutation of chromosome C_1 into chromosome C_1' and C_1''

V. EXPERIMENTAL RESULTS

A. Benchmark Test Instances

Two sets of instances are used to evaluate our modeling on existing algorithms. Each instance has 100 trips and is built on geographically realistic data [10]. All picking-up or delivery locations are geolocalized stops and the shortest duration matrix M is built using a GIS².

The first set, denoted '*Rnd100*', is composed of 10 randomly generated instances, whereas the instances of the second set, named '*Gravit100*', are generated by a *geographical* model of flows. The '*gravity model*' [14], as well as other interaction models, allows to quantify people flows in a geographical

²Geographical Information System

space according to a calibration based on inhabitant densities, the time dimension...

Set '*Rnd100*' has instances with an almost homogeneous distribution of customers while set '*Gravit100*' has instances with a non-homogeneous distribution representing flows between 8:00am and 9:00am. This effect results from the existence of attractive areas, which draw most flows instead of less attractive areas such as residential downtowns which are rather emission areas.

The algorithms the most suited to real-life applications can be highlighted by using realistic instances. Indeed, optimizing transport applications need efficient and robust algorithms.

B. Parameter Setting and Performance Assessment

The population of 100 individuals evolves over 10,000 generations. Crossover (x_r) and mutation (m_r) rates are respectively set to $x_r = 0.9$ and $m_r = 0.5$.

In addition, specific parameters exist for some algorithms. Hence, SPEA2 requires an internal archive whose size is fixed to 100 individuals. Following [13], the scaling factor κ of IBEA is set to 0.05.

A set of 20 runs per instance has been performed for each EA by using the ParadisEO-MOEO framework [15]. In order to evaluate the quality of the approximations for every instance we solved, we follow the protocol proposed in [16]. For a given instance, let Z^{all} denote the union of the outputs we obtained during all our experiments. We first compute a reference set Z_N^{\star} containing all the non-dominated points of Z^{all} . Second, we define $z^{min} = (z_1^{min}, \dots, z_n^{min})$ and $z^{max} = (z_1^{max}, \dots, z_n^{max})$, where z_k^{min} (resp. z_k^{max}) denotes the lower (resp. upper) bound of the k^{th} objective for all the points contained in Z^{all} . In order to give a roughly equal range to the objective functions, values are normalized with respect to z^{min} and z^{max} . Then, to measure the quality of an output set A in comparison to Z_N^{\star} , we compute the difference between these two sets by using the unary hypervolume metric [17], z^{max} being the reference point. The hypervolume difference indicator (I_H^-) computes the portion of the objective space that is dominated by Z_N^{\star} and not by A. Furthermore, we also consider the additive ϵ -indicator proposed in [17]. The unary additive ϵ -indicator ($I_{\epsilon+}^1$) gives the minimum factor by which an approximation A has to be translated in the objective space to dominate the reference set Z_N^{\star} . Note that both I_H^- and $I_{\epsilon+}^1$ -values are to be minimized.

Thus, for each test instance, we obtain $20 I_H^-$ measures and $20 I_{\epsilon+}$ measures corresponding to the 20 simulation runs *per* algorithm. Once all these values are computed, we perform a statistical analysis for a pairwise comparison of methods. To this end, we use the Wilcoxon signed rank test. For a given test instance, and with respect to a *p*-value of 0.05 and to the metric under consideration, this statistical test reveals if the sample of approximation sets obtained by a given search method is significantly better than the one of another search method, or if there is no significant difference between both of them.

C. Results and Discussion

Table I details a statistical comparison of each algorithm vs each other (symbols \prec , \succ , \equiv resp. indicate that the algorithm of a specific row is significantly 'better', 'worse' or 'equivalent' than the one of a specific column). The obtained results confirm that NSGA-II and IBEA are globally equivalent according to I_{H}^{-} , but that NSGA-II is always better (or at least equivalent) than IBEA according to $I_{\epsilon+}^{1}$. SPEA2 is always outperformed by the other methods. Thus, it appears that NSGA-II is better than both IBEA and SPEA2 for optimizing randomly generated instances of DARPTW.

Table II is the counterpart for the realistic instances. Contrary to Table I, IBEA is shown to be the best algorithm with respect to both I_H^- and $I_{\epsilon+}^1$ metrics (except for instance *Rnd100-4*). These unambiguous results indicate that IBEA is the most efficient (or at least equivalent) algorithm for optimizing realistic DARPTW instances while comparing pairwise algorithms. If we focus on the comparison of the two other algorithms, we see that SPEA2 is always better than NSGA-II with respect to I_H^- , while they are almost equivalent with respect to $I_{\epsilon+}^1$.

The main differences between the two kinds of instances are the densities of customers on the territory. Given the level of homogeneity of the distribution of the requests, the algorithms differently cope with the clustering effects. Hence, it seems that IBEA which uses a selection by indicator, appears to be more robust to the clustering effects due to the heterogeneous distribution of the customers.

VI. CONCLUSION AND PERSPECTIVES

This paper is devoted to solve a DARPTW applied to a Demand Responsive Transport service. After having reminded the general problem, an original evolutionary multi-objective approach was presented. Thanks to the Pareto-dominance principles, our approach is capable to build a set of distinct solutions for helping decision makers. To fit the evolutionary algorithms to be compared, a specific representation and variation operators were developed.

The candidate algorithms are: NSGA-II, IBEA and SPEA2. They were performed on two sets of transport instances. The first set is composed of random instances and the second of realistic instances. Such instances allow to assess the algorithm efficiency to cope with real-life applications.

The analysis of the results underline the more or less adapted behavior of the algorithms. It results of this comparative study that NSGA-II generally seems to be more efficient than IBEA for the random instances and SPEA2 is always outperformed by its counterparts. However, if we focus on the realistic instances, the results are opposite and bring to light that IBEA is the best algorithm for solving realistic DARPTW.

Given that the algorithms are sensitive to the distribution of customers and flows, a preemptive and relevant work would be to integrate the geographical dimension. Indeed, the better results obtained by IBEA on the realistic instances invite us to take this dimension into account. We believe that future research on more robust multiobjective evolutionary algorithms applied to real-life problems such as transportation will intensify. Moreover, the experiments on realistic instances indicate the algorithm sensibility to a specific topology. Then, it is necessary to evaluate the algorithms behavior according to the problem structures. Besides, an uncertainty integration [18] in the method will help us to develop dynamic DARPTW services.

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 TABLE I

 STATISTICAL COMPARISON FOR RANDOMLY GENERATED INSTANCES.

			r-			r 1	
		NII	I_H^- I	S2	NII	$\operatorname{I}^1_{\epsilon+}_{\operatorname{I}}$	S2
RO	NII	-	=	\succ	-	~	\succ
	I	≡	-	\succ	\prec	-	≡
	S2	\prec	\prec	-	ΥY	≡	-
R1	NII	-	\equiv	Υ		\succ	× ≺
	Ι	=	-	\succ	\prec	-	\succ
	S2	\prec	\prec	-	·ΥΥ	\prec	-
R2	NII	-	Ξ	$\stackrel{\scriptstyle \scriptstyle \times}{}$	-	Ξ	×
	Ι	=	-	\succ	\equiv	-	\succ
	S2	$ $ \prec	\prec	-	\parallel	\prec	-
R3	NII	-	\prec	\succ	-	\succ	\succ
	Ι	\succ	-	\succ	·ΥΥ	-	\succ
	S2	${\prec}$	\prec	-	\prec	\prec	-
R4	NII	-	≡	ΥΥ	-	\succ	\succ
	Ι	≡	-		\prec	-	\prec
	S2	$ $ \prec	\prec	-	·ΥΥ	<u>≻</u>	-
R5	NII	-	Ξ	\succ	י Y Y	\succ	\succ
	Ι	≡	-	\succ	\prec	-	=
	S2	\prec	\prec	-	\prec	Ξ	
R6	NII	-	=	\succ	-	\succ	\succ
	Ι	≡	-	\succ	\prec	-	\equiv
	S2	≺	\prec	-	·ΥΥ	\equiv	-
R7	NII	-	≡	$\stackrel{\scriptstyle \succ}{}$	• Y Y	\succ	\succ
	Ι	≡	-	\succ	\prec	-	\equiv
	S2	\prec	\prec	-	\prec	≡	-
R8	NII	-	Ξ	\prec	-	\succ	ΥY
	Ι	≡	-	\succ	• Y Y	-	\succ
	S2	\prec	\prec	-	\prec	\prec	-
R9	NII	-	Ξ	\succ	-	\succ	\succ
	I S2	∣≡ ≺	- ≺	\succ	- Y Y	-	\equiv
	S2	$ \prec$	\prec	-	\prec	\equiv	-

For each instance, the algorithms are opposed and compared by twos (NII: NSGA-II, I: IBEA, S2: SPEA2); Instance *Ri* indicates instance *Rnd100-i*

TABLE II

STATISTICAL COMPARISON FOR REALISTIC INSTANCES.

			I_H^- I			$I^1_{\epsilon+}$	
		NII	I	S2	NII	Ĭ	S2
G0	NII	-	\prec	\prec	-	\prec	Ξ
	Ι	\succ	-	\succ	\succ	-	\succ
	S2	\succ	\prec	-	≡	\prec	-
G1	NII	-	\prec	Ξ	-	\prec	Ξ
	Ι	\succ	-	\succ	\succ	-	\succ
	S2	≡	\prec	-	≡	\prec	-
G2	NII	-	\prec	\prec	-	\prec	\prec
	Ι	\succ	-	\succ	\succ	-	\succ
	S2	\succ	\prec	-	\succ	\prec	-
G3	NII	-	\prec	≡	-	\prec	Ξ
	Ι	≻	-	\succ	\succ	-	\succ
	S2	≡	\prec	-	\equiv	\prec	-
G4	NII	-	\prec	≡	-	Ξ	\succ
	Ι	≻	-	\succ	≡	-	\equiv
	S2	⊨	\prec	-	\prec	\equiv	-
G5	NII	-	\prec	\prec	-	\prec	≡
	Ι	\succ	-	\succ	\succ	-	\succ
	S2	\succ	\prec	-	\equiv	\prec	-
G6	NII	-	\prec	\prec	-	\prec	\prec
	Ι	\succ	-	\succ	\succ	-	\succ
	S2	\succ	\prec	-	\succ	\prec	-
G7	NII	-	\prec	\equiv	-	\prec	Ξ
	Ι	\succ	-	\succ	\succ	-	\succ
	S2	≡	\prec	-	\equiv	\prec	-
G8	NII	-	\prec	≡ ≻	-	\prec	=
	Ι	\succ	-	\succ	\succ	-	\succ
	S2	≡	\prec	-	\equiv	\prec	-
G9	NII	-	\prec	\prec	-	\prec	Ξ
	Ι	\succ	-	\succ	\succ	-	\succ
	S2	\succ	\prec	-	\equiv	\prec	-

For each instance, the algorithms are opposed and compared by twos (NII: NSGA-II, I: IBEA, S2: SPEA2); Instance *Gi* indicates instance *Gravit100-i*