

BiOWA for Preference Aggregation with Bipolar Scales: Application to Fair Optimization in Combinatorial Domains

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Abstract

We study the biOWA model for preference aggregation and multicriteria decision making from bipolar rating scales. A biOWA is an ordered doubly weighted averaging extending standard ordered weighted averaging (OWA) and allowing a finer control of the importance attached to positive and negative evaluations in the aggregation. After establishing some useful properties of biOWA to generate balanced Pareto-optimal solutions, we address fair biOWA-optimization problems in combinatorial domains. We first consider the use of biOWA in multi-winner elections for aggregating graded approval and disapproval judgements. Then we consider the use of biOWA for solving robust path problems with costs expressing gains and losses. A linearization of biOWA is proposed, allowing both problems to be solved by MIP. A path-ranking algorithm for biOWA optimization is also proposed. Numerical tests are provided to show the practical efficiency of our models.

1 Introduction

Decision making is often a matter of balancing pros and cons of possible choices. This is the case in multicriteria decision problems where the solutions are generally assessed w.r.t partially conflicting criteria. This also holds in collective decision making when agents have contrasted opinions about candidates. This happens too in decision under uncertainty where the utility of an action can vary drastically from one scenario to another. Very often, positive and negative perceptions co-exist in individual values and are represented using a bipolar utility scale in which 0 acts as a landmark separating the positive and negative sides [Dubois and Prade, 2008].

Yet, most normative decision methods use the same aggregation logic regardless the sign of outcomes under consideration. The utility scale is often treated as an interval scale and preferences are not impacted by positive affine transformations of the utility scale. For example, many aggregation functions $f : \mathbb{R}^n \rightarrow \mathbb{R}$ used in multicriteria analysis are such that $f(ax + b) = af(x) + b$ for all $x \in \mathbb{R}^n, a \in \mathbb{R}_+, b \in \mathbb{R}$. Hence if $f(x) > f(y)$ we also have $f(ax + b) = af(x) + b > af(y) + b = f(ay + b)$. In this case, 0 has no specific status.

It has been observed in different contexts that decision makers tend to think of outcomes relative to a certain reference point (often the status quo). They may exhibit different attitudes towards gains (i.e. outcomes above the reference point) and losses (i.e. outcomes below the reference point) and care generally more about negative outcomes than positive ones. In the field of decision under risk, this observation is at the origin of Prospect Theory [Kahneman and Tversky, 1979] and Cumulative Prospect Theory (CPT) [Tversky and Kahneman, 1992] that provide decision models able to incorporate the observed behaviors. Similarly, in the field of multicriteria analysis, extensions of various decision models have been proposed for handling bipolar scales and explaining observed preferences [Labreuche and Grabisch, 2006; Grabisch *et al.*, 2008; Grabisch *et al.*, 2009]. In particular, a generalization of the discrete Choquet integral is proposed for aggregating criterion values expressed on bipolar scales. The model is a kind of sophisticated weighted average that assigns different weights to criteria and coalitions of criteria depending on the nature of criterion values being positive or negative. This model formally includes the CPT model as special case [Labreuche and Grabisch, 2006].

In this paper we study a bipolar extension of the Ordered Weighted Averaging (OWA), with the aim of proposing new tools for fair optimization in combinatorial domains. The standard OWA operator is an aggregation function introduced in [Yager, 1998] formally defined, for all $x \in \mathbb{R}^n$, by the dot product $f_w(x) = w \cdot x_\uparrow$ where $w \in \mathbb{R}_+^n$ is a weighting vector and x_\uparrow is the vector obtained by rearranging the components of x in ascending order. It is therefore a weighted sum of the components x_i , each component representing the utility of solution x w.r.t a given point of view (e.g., a criterion, an agent, a scenario). However, we remark that the weights are not attached to criteria but to ranks of satisfaction due to the sorting of components. Function f_w is symmetric and weights only serve to control the importance attached to good or bad evaluations in the aggregation. OWA is often used with *decreasing weights* ($w_i \geq w_{i+1}, i = 1, \dots, n - 1$) in fair optimization because its maximization favors solutions with well-balanced profiles (a greater weight is assigned to lower components). More precisely, when w has decreasing components f_w is monotonically increasing w.r.t *Mean-Preserving Transfers Reducing Inequalities* (MPTRI), i.e., any transfer moving from $x = (x_1, \dots, x_n)$

3 BiOWA: an OWA for Bipolar Scales

A natural extension of OWA to bipolar scales, in the style of CPT, is the following;

Definition 1. Let $x \in \mathbb{R}^n$ be an evaluation vector and $\alpha, \beta \in \mathbb{R}_+^n$ two weighting vectors, the bipolar ordered weighted averaging (biOWA for short) is the aggregation function $g_{\alpha, \beta} : \mathbb{R}^n \rightarrow \mathbb{R}$ defined by:

$$g_{\alpha, \beta}(x) = \alpha \cdot x_{\uparrow}^+ - \beta \cdot x_{\downarrow}^- \quad (1)$$

where $x^+ = \max\{x, 0\}$, $x^- = \max\{-x, 0\}$ and x_{\uparrow} (resp. x_{\downarrow}) is a vector derived from x by rearranging its components in ascending (resp. descending) order.

We observe that, when $x \in \mathbb{R}_+^n$ then $x^+ = x$ and $x^- = 0$ therefore $g_{\alpha, \beta}(x) = f_{\alpha}(x)$. In this case biOWA reduces to a standard OWA. More generally, for any fixed $x \in \mathbb{R}^n$, there exists $w \in \mathbb{R}_+^n$ such that $g_{\alpha, \beta}(x) = f_w(x)$. The weighting vector w depends on x and is given by:

$$w_i = \begin{cases} \beta_i & \text{if } (x_{\uparrow})_i < 0 \\ \alpha_i & \text{otherwise} \end{cases} \quad i = 1, \dots, n \quad (2)$$

Hence any biOWA can be seen as a generalization of OWA using a non-constant weighting vector that may vary with x . Coming back to Example 2 we see that $g_{\alpha, \beta}$ used with $\alpha = (\frac{1}{2}, \frac{1}{2})$ and $\beta = (1, 0)$ describes the observed preferences:

$$\begin{aligned} g_{\alpha, \beta}(0, 5) &= \alpha \cdot (0, 5) - \beta \cdot (0, 0) = (\alpha_1, \alpha_2) \cdot (0, 5) = 2.5 \\ g_{\alpha, \beta}(2, 2) &= \alpha \cdot (2, 2) - \beta \cdot (0, 0) = (\alpha_1, \alpha_2) \cdot (2, 2) = 2 \\ g_{\alpha, \beta}(4, -4) &= \alpha \cdot (0, 4) - \beta \cdot (4, 0) = (\beta_1, \alpha_2) \cdot (-4, 4) = -2 \\ g_{\alpha, \beta}(-2, 1) &= \alpha \cdot (0, 1) - \beta \cdot (2, 0) = (\beta_1, \alpha_2) \cdot (-2, 1) = -1.5 \end{aligned}$$

Then, a natural question is whether biOWA is monotonic w.r.t weak Pareto-dominance. The answer is given below:

Proposition 1. Let $x, y \in \mathbb{R}^n$ such that $x_i \geq y_i$, for $i = 1, \dots, n$, one of these inequalities being strict. Let $\alpha, \beta \in \mathbb{R}_+^n$ be two weighting vectors with decreasing and strictly positive components, then $g_{\alpha, \beta}(x) > g_{\alpha, \beta}(y)$.

Proof. Since α and β have decreasing components, we have:

$$\alpha \cdot x_{\uparrow} = \min_{\pi \in \Pi} \sum_{i=1}^n \alpha_{\pi(i)} x_i \quad (3)$$

$$\beta \cdot x_{\downarrow} = \max_{\pi \in \Pi} \sum_{i=1}^n \beta_{\pi(i)} x_i \quad (4)$$

where Π is the set of all permutations defined on $(1, \dots, n)$. Moreover, since $x_i^+ \geq y_i^+$ for all i we have, for all $\pi \in \Pi$, $\alpha_{\pi(i)} x_i^+ \geq \alpha_{\pi(i)} y_i^+$ and therefore $\sum_{i=1}^n \alpha_{\pi(i)} x_i^+ \geq \sum_{i=1}^n \alpha_{\pi(i)} y_i^+$. Hence, taking the minimum over all $\pi \in \Pi$ we obtain $\alpha \cdot x_{\uparrow}^+ \geq \alpha \cdot y_{\uparrow}^+$ (a) by (3). Similarly, since $y_i^- \geq x_i^-$ for all i we have, for all $\pi \in \Pi$, $\beta_{\pi(i)} y_i^- \geq \beta_{\pi(i)} x_i^-$ and therefore $\sum_{i=1}^n \beta_{\pi(i)} y_i^- \geq \sum_{i=1}^n \beta_{\pi(i)} x_i^-$. Hence, taking the maximum over all $\pi \in \Pi$ we obtain $-\beta \cdot x_{\downarrow}^- \geq -\beta \cdot y_{\downarrow}^-$ (b) by (4). Moreover, $x_i > y_i$ holds for some i by hypothesis. Therefore at least one inequality (a) or (b) is strict. Then the result follows by summing (a) and (b) term by term. \square

A nice consequence of this proposition is that maximizing $g_{\alpha, \beta}$ with positive weights necessary leads to a Pareto-optimal solution. Yet, this does not mean that $g_{\alpha, \beta}$ -optimization favors well balanced solutions. Let us determine a set of necessary and sufficient conditions on (α, β) for $g_{\alpha, \beta}$ to be monotonic w.r.t MPTRI.

Proposition 2. $g_{\alpha, \beta}$ is monotonic w.r.t MPTRI if and only if α and β have decreasing components and $\beta_l \geq \alpha_{l+1}$ for all $l \in \{1, \dots, n-1\}$.

Proof. \Leftarrow We assume that α and β have decreasing components and that for all $l \in \{1, \dots, n-1\}$ $\alpha_l \geq \beta_{l+1}$. Let us consider any $x \in \mathbb{R}^n$ with $x_i < x_j$ for some pair (i, j) . Let $x' = [x_1, \dots, x_i + \lambda, \dots, x_j - \lambda, \dots, x_n]$ with $\lambda \in (0, x_j - x_i)$ the vector derived from x using a MPTRI. Let w and w' be the weighting vectors respectively derived from x and x' using Equation (2). If x_i and x_j are of the same sign, or if $\lambda > \max(-x_i, x_j)$ or if $\lambda < \min(-x_i, x_j)$, then x and x' have the same number of positive (resp. negative) components. Hence we have $w = w'$ and therefore $g_{\alpha, \beta}(x') = f_w(x')$. Moreover $f_w(x') \geq f_w(x) = g_{\alpha, \beta}(x)$ (the inequality is due to the fact that w has decreasing components). Therefore $g_{\alpha, \beta}(x') \geq g_{\alpha, \beta}(x)$. Now let us consider the case where x and x' have a different number of negative components (this occurs when $x_i < 0 < x_j$ and $\min(-x_i, x_j) \leq \lambda \leq \max(-x_i, x_j)$). This means that either component i or component j (not both) has a different sign when passing from x to x' . In the sequel we assume that the sign changes for component j (a symmetric proof could be done for the other case). Hence we can decompose the transfer into two successive ones: the first transfer decreases component j to 0 and the second transfer completes the first one by decreasing component j from 0 to $x_j - \lambda$. Let $x'' = (x_1, \dots, x_i + x_j, \dots, x_j - x_j, \dots, x_n)$ the vector we obtain after the first transfer and w'' the associated weighting vector using Equation (2). Then we have:

- Since $x''_j = 0$, component j is still non-negative after the first transfer, and then $w'' = w$. Hence we have $g_{\alpha, \beta}(x'') = f_w(x'')$. Moreover $f_w(x'') \geq f_w(x) = g_{\alpha, \beta}(x)$ (the inequality is due to the fact that w has decreasing components) and therefore $g_{\alpha, \beta}(x'') \geq g_{\alpha, \beta}(x)$.
- Since $x''_j = 0$, the component j does not impact the calculation of $g_{\alpha, \beta}(x'')$, which implies that $f_{w''}(x'') = f_{w'}(x'')$. Therefore we have $g_{\alpha, \beta}(x') = f_{w'}(x') \geq f_{w'}(x'') = g_{\alpha, \beta}(x'')$ (the inequality is due to the fact that w' has decreasing components).

Finally, we have $g_{\alpha, \beta}(x') \geq g_{\alpha, \beta}(x)$ in all cases. This shows that $g_{\alpha, \beta}$ is monotonic w.r.t MPTRI.

\Rightarrow Let x' be a solution obtained from x by a MPTRI. We can distinguish two possible cases where (α, β) do not fulfill the required conditions:

- Either α or β does not have decreasing components. In the former case we can consider a non-constant vector x with no negative component. Then $g_{\alpha, \beta}(x) = f_{\alpha}(x)$ is an OWA, known to be non-monotonic w.r.t MPTRI since components of α are not in the right order. The same reasoning applies to the latter case considering any other x with no positive component.
- $\exists l \in \{1, \dots, n-1\}$ with $\beta_l < \alpha_{l+1}$. Let us consider x such that $x_1 < \dots < x_l < 0 < x_{l+1} < \dots < x_n$ and x' the solution obtained from x by a MPTRI between x_{l+1} and x_l of size λ chosen sufficiently small

to preserve in x' the element's order of x . We have $g_{\alpha,\beta}(x') - g_{\alpha,\beta}(x) = \beta_l(x_l + \lambda) + \alpha_{l+1}(x_{l+1} - \lambda) - \beta_l x_l - \alpha_{l+1} x_{l+1} = \lambda(\beta_l - \alpha_{l+1}) < 0$. Then, $g_{\alpha,\beta}$ is not monotonic w.r.t MPTRI.

The conditions are therefore necessary and sufficient. \square

Proposition 2 characterizes the set of admissible parameters (α, β) for performing a fair optimization with a biOWA. Now, we study two applications of fair biOWA optimization, first in social choice and then in robust optimization.

4 Multi-Winner Election with Approval and Disapproval Ballots

Approval balloting is an evaluation system used in voting procedures in which each voter submits a ballot including the candidates he approves. It can be extended to richer systems involving approval and disapproval ballots [Felsenthal, 1989]. Such systems are often used in single-winner elections but can be extended to the multi-winner case [Kilgour, 2010]. We consider here a refinement of such voting systems where voters assign positive or negative grades to candidates to modulate approval and disapproval intensities.

More formally, let V be a set of n voters and C a set of m candidates. Let $v_{ij} \in \llbracket -T, T \rrbracket$ be the grade assigned to candidate j by voter i , T being the maximal admissible grade. We assume that preferences are additive over bundles and therefore the value of any subset $S \subseteq C$ for voter i is defined by $\sum_{j \in S} v_{ij}$. Assume we want to elect a committee of k candidates maximizing the voters' satisfaction, while paying a particular attention to unsatisfied voters (with negative satisfactions). We propose to use a biOWA to model the overall value of any subset. Hence we can formulate the biOWA Multi-Winner Election Problem (biOWA-MWE Problem) with the following mixed integer program:

$$\begin{aligned} & \max g_{\alpha,\beta}(x_1, \dots, x_n) \\ \text{s.t. } & \begin{cases} x_i = \sum_{j=1}^m v_{ij} y_j & i = 1, \dots, n \\ \sum_{j=1}^m y_j = k \\ y_j \in \{0, 1\}, j = 1, \dots, m \end{cases} \end{aligned} \quad (5)$$

where y_j is the decision variable relative to the selection of candidate j , $j = 1, \dots, m$. For any fixed k , the biOWA-MWE problem is polynomial. Indeed, we can enumerate the $\binom{m}{k}$ subsets of size k and return the $g_{\alpha,\beta}$ -optimal subset. The general problem is NP-hard because biOWA optimization includes min-max optimization and the min-max version of the problem, a.k.a *the minimum selecting items problem*, is known to be NP-hard for two scenarios and strongly NP-hard for an unbounded number of scenarios [Kasperski and Zieliński, 2015]. In order to solve the biOWA-MWE problem using mixed-integer programming, we establish a useful result for linearizing $g_{\alpha,\beta}$:

Proposition 3. For all $x \in \mathbb{R}^n$, and all $\alpha, \beta \in \mathbb{R}_+^n$ two weighting vectors having decreasing components we have:

$$g_{\alpha,\beta}(x) = \min_{\tau \in \Pi} \sum_{i=1}^n (\alpha_{\tau(i)} x_i^+ - \beta_{\tau(i)} x_i^-) \quad (6)$$

with Π the set of permutations on $\{1, \dots, n\}$.

Proof. Due to Equations (3) and (4) we have:

$$\begin{aligned} g_{\alpha,\beta}(x) &= \min_{\pi \in \Pi} \sum_{i=1}^n \alpha_{\pi(i)} x_i^+ - \max_{\pi' \in \Pi} \sum_{i=1}^n \beta_{\pi'(i)} x_i^- \\ &= \min_{\pi \in \Pi} \sum_{i=1}^n \alpha_{\pi(i)} x_i^+ + \min_{\pi' \in \Pi} \sum_{i=1}^n \beta_{\pi'(i)} (-x_i^-) \\ &\leq \min_{\tau \in \Pi} \sum_{i=1}^n (\alpha_{\tau(i)} x_i^+ - \beta_{\tau(i)} x_i^-) \end{aligned}$$

Now we establish the reverse inequality. Let π_* a permutation such that $\alpha \cdot x^+ = \sum_{i=1}^n \alpha_{\pi_*(i)} x_i^+$ and π'_* a permutation such that $\beta \cdot x^- = \sum_{i=1}^n \beta_{\pi'_*(i)} x_i^-$. Now, let us consider any permutation τ_* of $(1, \dots, n)$ such that $\tau_*(i) = \pi_*(i)$ if $x_i > 0$ and $\tau_*(i) = \pi'_*(i)$ if $x_i < 0$, $\tau_*(i)$ being chosen arbitrarily for all i such that $x_i = 0$ to complete the permutation. By construction we have:

$$\begin{aligned} \min_{\tau \in \Pi} \sum_{i=1}^n (\alpha_{\tau(i)} x_i^+ - \beta_{\tau(i)} x_i^-) &\leq \sum_{i=1}^n (\alpha_{\tau_*(i)} x_i^+ - \beta_{\tau_*(i)} x_i^-) \\ &= \sum_{i=1}^n \alpha_{\pi_*(i)} x_i^+ - \sum_{i=1}^n \beta_{\pi'_*(i)} x_i^- = g_{\alpha,\beta}(x). \quad \square \end{aligned}$$

Proposition 3 shows that $g_{\alpha,\beta}(x)$ can be computed, for any fixed $x \in \mathbb{R}^n$, by solving the following linear program:

$$\begin{aligned} & \min \sum_{i=1}^n \sum_{j=1}^n (\alpha_i x_j^+ - \beta_j x_j^-) p_{ij} \\ \text{s.t. } & \begin{cases} \sum_{i=1}^n p_{ij} = 1 & j = 1, \dots, n \\ \sum_{j=1}^n p_{ij} = 1 & i = 1, \dots, n \\ p_{ij} \geq 0, i, j = 1, \dots, n \end{cases} \end{aligned} \quad (7)$$

In the above program, variables p_{ij} are used to model permutations of $(1, \dots, n)$. Let us first assume that p_{ij} are boolean variables. In this case they are able to model any permutation $\pi \in \Pi$ by setting $p_{ij} = 1$ if $\pi(i) = j$ and 0 otherwise. Now, if we relax variables p_{ij} , these permutations are represented by the vertices of the convex polyhedron defined by variables p_{ij} (vertices are integral). Hence, due to the linearity of the objective function w.r.t variables p_{ij} , one can relax the integrity of variable p_{ij} without changing the optimal value of the problem. This argument is used in [Chassein and Goerigk, 2015] to model permutations of OWA and still applies to permutations used in biOWA. The dual formulation of the above linear program (7) reads as follows:

$$\begin{aligned} & \max \sum_{i=1}^n (n_i + p_i) \\ \text{s.t. } & n_i + p_j \leq \alpha_i x_j^+ - \beta_j x_j^- \quad i, j = 1, \dots, n \\ & n_i, p_i \in \mathbb{R}, i = 1, \dots, n \end{aligned} \quad (8)$$

Using this formulation, we propose a mixed integer program (\mathcal{P}) to maximize $g_{\alpha,\beta}(x)$ over a set X :

$$(\mathcal{P}) \quad \begin{cases} \max \sum_{i=1}^n (n_i + p_i) \\ n_i + p_j \leq \alpha_i x_j^+ - \beta_i x_j^- & i, j = 1, \dots, n \\ x_i = x_i^+ - x_i^- & i = 1, \dots, n \\ 0 \leq x_i^+ \leq t_i \times M & i = 1, \dots, n \\ 0 \leq x_i^- \leq (1 - t_i) \times M & i = 1, \dots, n \\ x \in X \\ x_i, x_i^+, x_i^- \geq 0, i = 1, \dots, n \\ n_i, p_i \in \mathbb{R}, i = 1, \dots, n \\ t_i \in \{0, 1\}, i = 1, \dots, n \end{cases}$$

The integer variables $t_i, i = 1, \dots, n$ are used to decide whether x_i is positive or not. The M constant is used as usual to model disjunctive constraints depending on the sign of x_i .

Program \mathcal{P} for biOWA optimization can be specialized to solve the biOWA-MWE problem. It is sufficient to insert m boolean variables y_j modeling elementary decisions on candidates ($y_j = 1$ iff the candidate is selected in the current solution). Then $x \in X$ must be replaced by the constraints $x_i = \sum_{j=1}^m v_{ij} y_j, i = 1, \dots, n$ and $\sum_{j=1}^m y_j = k$. We obtain a mixed-integer program with $n + m$ boolean variables, $5n$ real variables and $n^2 + 4n + 1$ linear constraints. Note that proposition 3 enables to reduce both the number of variables and constraints in this program. We indeed used only one set of permutation variables instead of the two needed if biOWA were only seen as a difference of two OWAs.

We have implemented the above model using the Gurobi 7.5.2 solver on a computer with 12GB of RAM, a Intel(R) Core(TM) i7 CPU 950 @ 3.07GHz processor. We used instances of the BiOWA-MWE problem of different sizes, the number of candidates (m) ranging from 20 to 100 and the number of voters (n) ranging from 10 to 100. Votes v_{ij} are randomly generated in the range $\llbracket -10, 10 \rrbracket$ and $k = \frac{m}{2}$. Vectors α and β are randomly drawn and satisfy the conditions of Proposition 2. For each pair (n, m) , the average time given in Table 2 is expressed in seconds and computed over 20 runs, with a timeout set to 1200 seconds.

Another linearization of biOWA could be obtained by exploiting a linearization of OWA due to Ogryczak [Ogryczak, 2003]. We have implemented and tested this second option but it appears to be less efficient. Its presentation has been omitted here to save space.

5 Robust Paths with Gains and Losses

The ability of OWA or biOWA optimization to generate solutions having a balanced utility vector can also be exploited in the context of robust optimization. The need of robustness appears in decision problems under uncertainty,

n	$m = 20$	$m = 50$	$m = 100$
10	0.04	0.06	0.18
20	0.13	0.21	0.53
50	0.79	1.48	9.96
100	5.13	25.78	342.24

Table 2: Times (s) obtained by MIP for the biOWA-MWE

when a number of plausible scenarios must be considered, each scenario impacting differently the values of solutions. In such contexts, uncertainty aversion leads to prefer robust solutions, i.e., solutions having a balanced profile over scenarios. In this context, the use of OWA-optimization has been closely investigated [Perny and Spanjaard, 2003; Kasperski and Zielinski, 2015]; It is therefore natural to study the use of biOWA when positive and negative values are present. For example, let us consider the robust path problem on a graph valued in a bipolar scale, modeling gains or losses induced by transitions between nodes.

More formally, let $G = (V, E)$ be an acyclic directed graph where V is a finite set of nodes and E the set of arcs representing possible transitions between nodes. For all $i = 1, \dots, n$, let $u_i : E \rightarrow \mathbb{R}$ be a valuation function defined on E , $u_i(e)$ representing the gain (or loss) attached to e in scenario i . Valuations of arcs are supposed to be additive along a path and we want to find a biOWA-optimal path from s to t in G . This problem is NP-hard because biOWA includes OWA as special case and the OWA optimal path is known to be NP-hard for 2 scenarios and strongly NP-hard for an unbounded number of scenarios [Kasperski and Zielinski, 2015].

Program \mathcal{P} introduced in Section 4 can be specialized to determine a biOWA-optimal path problem using standard max flow formulations of paths problems. To define the feasible set of solutions X let us introduce $I(v)$ (resp. $O(v)$) the set of edges entering in v (resp. leaving v) for any $v \in V$ and y_e a boolean variable (set to 1 if e is selected in the path), for all $e \in E$. Then we add the following constraints:

$$\begin{aligned} x_i &= \sum_{e \in E} u_i(e) y_e, \quad i = 1, \dots, n \\ \sum_{e \in O(k)} y_e - \sum_{u \in I(k)} y_u &= \begin{cases} 1 & \text{if } k = s \\ -1 & \text{if } k = t \\ 0 & \text{if } k \in V \setminus \{s, t\} \end{cases} \end{aligned}$$

We implemented the above model using the Gurobi solver on the computer described in Section 4. We made tests on random instances including up to $m = 4000$ nodes and $n = 7$ scenarios (in robust paths problems with a finite set of scenarios, one generally considers only a small number of scenarios, but 2 are sufficient to obtain a NP-hard problem). To generate instances, nodes are uniformly distributed on 25 different layers and arcs are randomly generated between nodes of consecutive layers, with a probability $\frac{1}{2}$. Valuations of arcs are generated within $\llbracket -100, 100 \rrbracket$ with an uniform distribution. Weights α and β are randomly generated and satisfy the conditions of Proposition 2. Table 3 gives the average computational times obtained over 20 runs, with a timeout set to 1200 seconds for each run.

We can see that this new specialization of program \mathcal{P} intro-

m	$n = 3$	$n = 5$	$n = 7$
500	1.08	1.19	2.81
1000	3.86	4.27	15.89
2000	47.91	56.43	173.49
3000	33.15	61.88	374.66
4000	50.98	103.01	444.50

Table 3: Times (s) obtained by MIP for biOWA-optimal paths

duced in Section 4 can solve all instances in very reasonable time. To go further on the biOWA optimal path problem, we also investigated combinatorial algorithms in graphs. First, it is important to remark that it is not possible to construct a biOWA-optimal path with a simple dynamic programming algorithm progressively extending biOWA-optimal subpaths. This is due to the fact that preferences induced by a biOWA do not satisfy the Bellman principle. This phenomenon is already known for OWA and can be illustrated by the following:

Example 3. Consider an instance $G = (V, E)$ with $V = \{1, 2, 3, 4, 5\}$ and $E = \{(1, 2), (1, 3), (2, 4), (3, 4), (4, 5)\}$ and two scenarios leading to the following outcomes:

arcs	(1, 2)	(1, 3)	(2, 4)	(3, 4)	(4, 5)
outcomes	(6, -2)	(3, -1)	(-1, 2)	(-1, 4)	(0, 5)

We look for the $g_{\alpha, \beta}$ -maximal path from 1 to 5 with $\alpha = (\frac{2}{3}, \frac{1}{3})$ and $\beta = (\frac{3}{4}, \frac{1}{4})$. Subpath 1-2-4 leads to outcomes (5, 0) and $g_{\alpha, \beta}(5, 0) = \frac{5}{3}$. Subpath 1-3-4 leads to outcomes (2, 3) and $g_{\alpha, \beta}(2, 3) = \frac{7}{3}$. Therefore 1-3-4 is the best path from 1 to 4 according to biOWA valuations. Moreover, 1-3-4-5 yields (2, 3) + (0, 5) = (2, 8) and $g_{\alpha, \beta}(2, 8) = \frac{10}{3}$ while 1-2-4-5 yields (5, 0) + (0, 5) = (5, 5) and $g_{\alpha, \beta}(5, 5) = 5$. Hence the biOWA-optimal path from 1 to 5 is 1-2-4-5 although the subpath 1-2-4 was not optimal from 1 to 4. This violation of the Bellman principle prevents local pruning of subpaths based on $g_{\alpha, \beta}$ values in a constructive algorithm.

To overcome the problem we use a path-ranking algorithmic scheme which has been successfully used to solve multiobjective optimization problems in graphs under nonlinear scalarization models [Galand and Perny, 2007]. Let g be a non-linear function defining the scalar value of a path from its outcome vector, and assume that we can approximate g by a linear function h such that, for all $x \in \mathbb{R}^n$ verifies $g(x) \leq h(x)$. Then the path-ranking scheme consists in 1) scalarizing outcome vectors attached to arcs using the h function and 2) launching the enumeration of paths by decreasing values of h . To implement this approach, we need an efficient path-ranking algorithm. We use here an improved version of Eppstein algorithm proposed in [Jiménez and Marzal, 2003].

At any step of the ranking process, let x be the current solution and x^* the solution with the highest g -value among those obtained so far. Whenever $g(x^*) \geq h(x)$, we know that x^* is the optimal solution for this instance. Indeed, we have, for any y that will be enumerated after x in the ranking algorithm, $g(x^*) \geq h(x) \geq h(y) \geq g(y)$. Hence y cannot improve x^* and the ranking process can be stopped, x^* is the g -optimal solution.

To apply this scheme to $g = g_{\alpha, \beta}$ we need a proper function h . To this end, we establish the following inequality:

Proposition 4. Let α and β be two positive vectors having decreasing components such that $\sum_{i=1}^n \alpha_i = \sum_{i=1}^n \beta_i = 1$, then for all $x \in \mathbb{R}^n$ we have: $g_{\alpha, \beta}(x) \leq \frac{1}{n} \sum_{i=1}^n x_i$.

Proof. It is well known that: $\alpha \cdot x_{\uparrow} \leq \frac{1}{n} \sum_{i=1}^n x_i$ when α has decreasing components. As a direct consequence, we also have: $\beta \cdot x_{\downarrow} \geq \frac{1}{n} \sum_{i=1}^n x_i$ when β has decreasing components. Hence we have: $g_{\alpha, \beta} = \alpha \cdot x_{\uparrow} - \beta \cdot x_{\downarrow} \leq \frac{1}{n} \sum_{i=1}^n x_i^+ - \frac{1}{n} \sum_{i=1}^n x_i^- = \frac{1}{n} \sum_{i=1}^n (x_i^+ - x_i^-) = \frac{1}{n} \sum_{i=1}^n x_i$ \square

m	$n = 3$	$n = 5$	$n = 7$
500	0.16	2.99	38.14
1000	0.52	3.72	78.90
2000	1.92	10.10	367.58
3000	3.91	9.53	221.71
4000	6.85	15.40	352.4

Table 4: Times (s) obtained by ranking for biOWA-optimal paths

Thus, the average can be used as a valid h function in the ranking scheme, provided we use normalized vectors α and β . Table 4 provides the results obtained with the path-ranking algorithm for the biOWA-optimal path problem. The instances are generated as before, using layered graphs. Computational times are obtained by averaging over 20 runs, with a timeout set to 1200 second for each run.

If we compare the results given in Table 4 with previous results obtained using Gurobi and the MIP model (Table 3), we observe that the ranking-path algorithm is faster on average. A deeper analysis shows that this is due to the fact that the $g_{\alpha, \beta}$ -optimal path is often very well ranked according to the average (h). On the other hand, there exist instances where the $g_{\alpha, \beta}$ -optimal path appears much later in the ranking process, yielding worse computational times. The previous approach based on MIP is less sensitive to this phenomenon.

6 Conclusion

In this paper we have identified the conditions under which a biOWA is monotonic w.r.t mean-preserving transfers reducing inequalities. This justifies its use in multiobjective optimization contexts when one wants to favor the determination of *balanced* Pareto-optimal solutions. Then we have proposed and tested computational models allowing to effectively determine biOWA-optimal solutions on combinatorial domains. We presented two possible applications, first in the context of fair multiagent optimization and then in the context of robust optimization. Many other problems could be considered as well due to generality of the proposed computational models (MIP and ranking algorithm).

One specificity of biOWA is to perform a symmetric aggregation of its arguments. Although this property is highly desirable in Social Choice (when every agent has the same importance) or in robust optimization under total uncertainty (when all scenarios are equally likely), there are other situations where different weights or probabilities must be attached to components (criteria, scenarios or individual values), thus breaking the symmetry. The computational models proposed in this paper could very likely be extended to the case of weighted OWA [Torra, 1997], CPT and other classes of general Choquet integrals [Grabisch *et al.*, 2009] to perform weighted aggregation on bipolar valuation scales in optimization problems.

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