

A fast dynamic programming multi-objective knapsack problem

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Abstract

This work addresses... The Multidimensional Objective knapsack programming.
The dynamic programming method... The data structure...

1 Introduction

2 The Multiobjective Knapsack Problem

A general multiobjective optimization problem can be described as a vector function f that maps a tuple of n parameters (decision variables) to a tuple of m objectives. Formally:

$$\begin{aligned} \min/\max \mathbf{y} &= f(\mathbf{x}) = (f_1(\mathbf{x}), f_2(\mathbf{x}), \dots, f_m(\mathbf{x})) \\ \text{subject to } \mathbf{x} &= (x_1, x_2, \dots, x_n) \in X \end{aligned}$$

where \mathbf{x} is the *decision vector*, X denotes the set of feasible solutions, and \mathbf{y} is the *objective vector* where each objective has to be minimized (or maximized).

Considering two decision vectors $\mathbf{a}, \mathbf{b} \in X$, \mathbf{a} is said to *dominate* \mathbf{b} if, and only if \mathbf{a} is at least as good as \mathbf{b} in all objectives and better than \mathbf{b} in at least one objective. For shortening we will say that \mathbf{a} dominates \mathbf{b} by saying $dom(\mathbf{a}, \mathbf{b})$. Formally:

$$dom(\mathbf{a}, \mathbf{b}) = \begin{cases} \forall i \in \{1, 2, \dots, m\} : f_i(\mathbf{a}) \geq f_i(\mathbf{b}) \text{ and} \\ \exists j \in \{1, 2, \dots, m\} : f_j(\mathbf{a}) > f_j(\mathbf{b}) \end{cases}$$

A feasible solution $\mathbf{a} \in X$ is called *efficient* if its not dominated by any other feasible solution. The set of all efficient solutions of a multiobjective optimization problem is known as *Pareto optimal*. Solving a multiobjective problem consists in giving its Pareto optimal set.

An instance of a multiobjective knapsack problem (MOKP) with m objectives consists of an integer capacity $W > 0$ and n items. Each item i has a

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positive weight w_i and non negative integer profits $p_i^1, p_i^2, \dots, p_i^m$. Each profit p_i^k represents the contribution of the i -th item for k -th objective. A solution is represented by a set $\mathbf{s} \subseteq \{1, \dots, n\}$ containing the indexes of the items included in the solution. A solution is feasible if the total weight included in the knapsack does not exceed its capacity. Formally the definition of the problem is:

$$\begin{aligned} \max f(\mathbf{s}) &= (f_1(\mathbf{s}), f_2(\mathbf{s}), \dots, f_m(\mathbf{s})) \\ \text{subject to } w(\mathbf{s}) &< W \\ \mathbf{s} &\subseteq \{1, \dots, n\} \end{aligned}$$

where

$$\begin{aligned} f_j(\mathbf{s}) &= \sum_{i \in \mathbf{s}} p_i^j \\ w(\mathbf{s}) &= \sum_{i \in \mathbf{s}} w_i \end{aligned}$$

The MOKP is considered a \mathcal{NP} -Hard problem since it is a generalization of the well-known 0–1 knapsack problem, in which $m = 1$. It is quite difficult to determine the Pareto optimal set for the MOKP, especially for high dimension instances, in which the solution it self (Pareto optimal set) tends to grow exponentially. Even for the bi-objective case, small problems may prove intractable. For this reason we are interested in developing efficient methods for handling large solution sets, which may bring tractability to previously intractable instances.

3 The Dynamic Programing Algorithm

Paragrafo de introducao da secao, justificando toda a explicacao que segue..

This dynamic programming algorithm is based on the classical Nemhauser-Ullmann (NU) algorithm proposed in [?] which consists of a sequential process of n stages. At each stage a new set of solution is constructed using a concept of domination for knapsack problems proposed by Weingartner and Ness [?]. At the the n -th stage the Pareto optimal set obtained.

Some property of the MOKP will be explored three filter as optimizations for the trying to reduce the number of solutions handled on mid stages of the algorithm. We will introduce the dynamic programming (DP) algorithm for the MOKP proposed in [?].

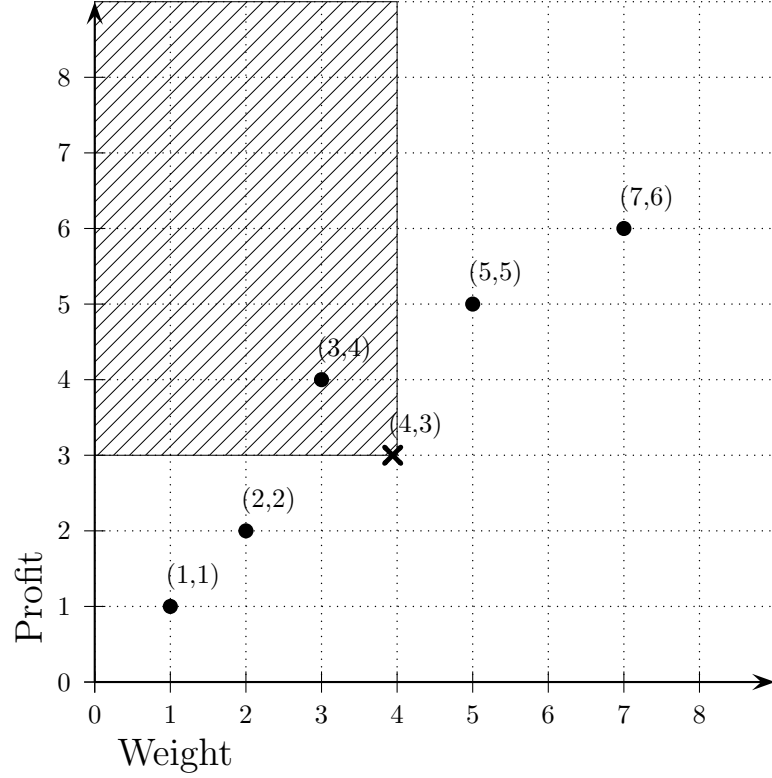
Before we will present some definitions...

3.1 The NU algorithm – the knapsack dominance

A solution \mathbf{x} is called *deficient* if it has available space to fit one or more item, i.e., $w(\mathbf{x}) + \min\{w_i : i \notin \mathbf{x}\} \leq W$. Considering two solution $\mathbf{x}, \mathbf{y} \subseteq \{1, \dots, n\}$ we say \mathbf{x} *knapsack-dominates* \mathbf{y} if it dominates \mathbf{y} and does not weight more than \mathbf{y} . For shortening we will say that \mathbf{x} knapsack-dominates \mathbf{y} by saying

$dom_k(\mathbf{x}, \mathbf{y})$. Formally:

$$dom_k(\mathbf{x}, \mathbf{y}) = \begin{cases} dom(\mathbf{x}, \mathbf{y}) & \text{and} \\ w(\mathbf{x}) \leq w(\mathbf{y}) \end{cases}$$



The pseudocode for the algorithm is presented in Algorithm 1. At the k -th stage the algorithm receives a set S^{k-1} of solutions and generates the set S^k of solutions that correspond to subsets containing exclusively the first k items, i.e., $\forall \mathbf{x} \in S^k, \mathbf{x} \subseteq \{1, \dots, k\}$.

This is done by expanding S^{k-1} by adding a copy of each solution with the inclusion of k item (line 4). We will refer as *partial solutions* all the solutions handled by stages prior to n -th stage.

The clever part of the algorithm is that it uses the concept of knapsack dominance to filter solutions that will not lead to efficient solutions (line 5). Considering two partial solutions $\mathbf{x}, \mathbf{y} \in S^k$, if \mathbf{x} is knapsack-dominated by \mathbf{y} then we may discard \mathbf{x} since all solutions generated from \mathbf{x} will be dominated by those generated from \mathbf{y} .

Algorithm 1 Basic dynamic programming algorithm for MOKP

```
1: function DP( $\mathbf{p}, \mathbf{w}, W$ )
2:    $S^0 = \{\emptyset\}$ 
3:   for  $k \leftarrow 1, n$  do
4:      $S_*^k = S^{k-1} \cup \{\mathbf{x} \cup k \mid \mathbf{x} \in S^{k-1}\}$   $\triangleright$  solutions extension
5:      $S^k = \{\mathbf{x} \mid \nexists \mathbf{a} \in S_*^k : \text{dom}_k(\mathbf{a}, \mathbf{x})\}$   $\triangleright$  partial dominance filter
6:   end for
7:    $P = \{\mathbf{x} \mid \nexists \mathbf{a} \in S^n : \text{dom}(\mathbf{a}, \mathbf{x})\}$   $\triangleright$  dominance filter
8:   return  $P$ 
9: end function
```

3.2 Avoiding deficient solutions

The first optimization that can be made on Algorithm 1 is avoiding the generation of deficient solutions. At k -th stage all previous solution is copied to the new solution set without adding k -th item (line 4). However preserving solutions with a lot of space left, concerning the remaining items, may lead to deficient solutions.

Theorem 1. *Considering the k -th stage of the algorithm and $\mathbf{x} \in S^{k-1}$. If $w(\mathbf{x}) + \sum_{i \in \{k, \dots, n\}} w_i \leq W$ than*

Considering the k -th stage, if a partial solution $\mathbf{x} \in S^{k-1}$ has enough space to fit all remaining items, i.e., $w(\mathbf{x}) + \sum_{i=k}^n w_i \leq W$, \mathbf{x} may be discarded and only $\mathbf{x} \cup \{k\}$ kepted, once keeping \mathbf{x} will certainly lead to deficient solutions.

3.3 Removing unpromissing solutions

Another optimization that can be applied on later stages is filtering unpromissing solutions by computing upper bounds for its objectives functions and comparing it with the set of available lower bounds. An upper(lower) bound of a solution is an upper(lower) limit each objective value can achieve given its remaining capacity and the remaining items. If the upper bound of a solution is dominated by an existing lower bound, that solution can be discarded since it will not generate an efficient solution.

A lower bound of a solution can be easily computed...
Formally:

$$lb(\mathbf{x}) = (lb_1(\mathbf{x}), \dots, lb_m(\mathbf{x}))$$

where

$$lb_i(\mathbf{x}) = \dots$$

Algorithm 2 Bazgan’s DP algorithm for the MOKP

```
1: function BAZDP( $\mathbf{p}, \mathbf{w}, W$ )
2:    $S^0 = \{\emptyset\}$ 
3:   for  $k \leftarrow 1, n$  do
4:     2...
5:   end for
6:    $P = \{\mathbf{x} \mid \nexists \mathbf{a} \in S^n : \text{dom}(\mathbf{a}, \mathbf{x})\}$  ▷ dominance filter
7:   return  $P$ 
8: end function
```

3.4 Item order

4 The use of k -d tree

The k -d tree is a type of binary search tree for indexing multidimensional data with simple construction and low space usage. Despite its simplicity it efficiently supports operations like nearest neighbour search and range search [?]. For those reasons k -d tree is widely used on spacial geometry algorithms [?, ?], clustering [?, ?] and graphic rendering algorithms [?].

Like a standard binary search tree, the k -d tree subdivides data at each recursive level of the tree. Unlike a standard binary tree, that users only one key for all levels of the tree, the k -d tree uses k keys and cycles through these keys for successive levels of the tree.

Concerning it’s efficiency, it is important to consider the number of dimensions k -d tree is indexing. As a general rule, a k -d tree is suitable for efficiently indexing of n elements if n is much greater than 2^k . Otherwise, when k -d tree are used with high-dimensional data, most of the elements in the tree will be evaluated and the efficiency is no better than exhaustive search [?].

Indexing the solutions and range operations.

Tends to increase the feasibility on problems with higher dimensions.

5 Computational experiments

- Base de dados utilizaca
- Parametros dos algoritmos
- Anlise dos resultados (comparao)

6 Conclusions and future remarks

- Conclues dos resultados
- Trabalhos futuros

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