



**Design of approximation algorithms pdf** 



Scheduling to Minimize Total Weighted Completion Time Performance Guarantees of LP-Based Heuristics and Lower Bound

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Abstract. There has been recent success in using polyhedral formula-tions of scheduling problems not only to obtain good lower bounds in practice but also to develop provably good approximation algorithms. Most of these formulations rely on binary decision variables that are a kind of assignment variables. We present quite simple polynomial-time approximation algorithms that are based on linear programming formulations with completion time variables and give the best known performance guarantees for minimizing the total weighted completion time in several scheduling environments. This amplifies the importance of (appropriate) polyhedral formulations in the design of approximation algorithms with good worst-case performance guarantees. In particular, for the problem of minimizing the total weighted com-pletion time on a single machine subject to precedence constraints we present a polynomial-time approximation algorithm with performance ratio better than 2. This outperforms a  $(4+\epsilon)$ -approximation algorithm very recently proposed by Hall, Shnoys, and Wein that is based on time-indexed formulations. A slightly extended formulation leads to a performance guarantee of 3 for the same problem but with release dates. This improves a factor of 5.83 for the same problem and even the 4-approximation algorithm for the problem with release dates but without precedence constraints, both also due to Hall, Shnoys, and Wein. By introducing new linear inequalities, we also show how to extend our technique to parallel machines. Finall, Shunoys, and Wein. By introducing new linear inequalities, we also show how to extend our technique to parallel machines. Finally, for the flow shop problem to minimize the total weighted completion time with both precedence constraints and release dates we present the first approximation algorithm that achieves a worst-case performance guarantee that is linear in the number of machines. We even extend this to multiprocessor flow shop scheduling. The proofs of these results also imply guarant

## Algorithms for Generating Pareto fronts of Multi-objective Integer and Mixed-Integer **Programming Problems**

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Abstract Multi-objective integer or mixed-integer programming problems typically have disconnected feasible domains, making the task of constructing an approximation of the Pareto front challenging. The present paper shows that certain algorithms which were originally devised for continuous problems can be successfully adapted to approximate the Pareto front for integer, and mixed-integer, multi-objective optimization problems. Relationships amongst various scalarization techniques are established to motivate the choice of a particular scalarization in these algorithms. The proposed algorithms are tested by means of two-, three- and four-objective integer and mixed-integer problems, and comparisons are made. In particular, a new four-objective algorithm is used to solve a rocket injector design problem with a discrete variable, which is a challenging mixed-integer programming problem.

Key words: Multi-objective optimization, Integer programming, Mixed-integer programming, Scalarization, Pareto front, Efficient set, Numerical methods.

AMS subject classifications. 90C26, 90C29, 90C30, 90C56.

## 1 Introduction

Multi-objective optimization is concerned with simultaneous minimization of several conflicting objective functions. The aim is to obtain/approximate, and view, the set of trade-off, or compromise, solutions. This set of trade-off solutions is referred to, in the current article, as the (weak) Pareto front, or the (weak) efficient set, of the problem-a more precise definition is to be provided in Section 2. Once the set of Pareto points is at hand, the decision maker conveniently selects a suitable solution from this set, usually based on an additional criterion. These kinds of problems arise in many multidisciplinary applications, for example, in health [9, 52, 61], engineering [30, 50], mining [26, 59], and finance [45, 49].

If all variables of a multi-objective optimization problem belong to a continuous set, then the problem is called a multi-objective continuous optimization problem. If, on the other hand, the variables belong to a subset of integers, then the problem is a multi-objective integer programming one. When the model involves both integer and continuous variables the problem is referred to as a multi-objective mixed-integer programming problem. The latter case arises in many real-life applications, for example, the knapsack [10,28] and shortest-path problems [13,54]. Other types of mixed-integer multi-objective problems can be found in [43].

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## The degenerate primer design problem

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ABSTRACT A PCR primer sequence is called degenerate if some of ts positions have several possible bases. The degeneracy of the primer is the number of unique sequence combina tions it contains. We study the problem of designing a pair of primers with prescribed degeneracy that match a maximum number of given input sequences. Such problems occur when studying a family of genes that is known only in part, or is known in a related species. We prove that We may have to compromise by aiming to match many various simplified versions of the problem are hard, show the polynomiality of some restricted cases, and develop develop an ad hoc method for designing primers that will approximation algorithms for one variant. Based on these algorithms, we implemented a program called HYDEN for designing highly-degenerate primers for a set of genomic objective Degenerate Primer Design (DPD). nces. We report on the success of the program in an experimental scheme for identifying all human offactory ceptor (OR) genes. In that project, HYDEN was used to design primers with degeneracies up to 1010 that amplified with high specificity many novel genes of that family, tripling the number of OR genes known at the time. Availability: Available on request from the authors. Contact: chaiml@tau.ac.il; rshamir@tau.ac.il Keywords: PCR primer design; degenerate primers; optiization; human offactory subgenome; protein families. INTRODUCTION

## the sequences without gaps, count the number of differen nucleotides in each position along the alignment and seek a primer-length window (typically 20-30) where the product of the counts is low. Such a solution is insufficien because of gaps, the inappropriate objective function of the alignment, and, most notably, the exceedingly high degeneracy: When degeneracy is too high, unrelated equences may be amplified as well, losing specificity but not necessarily all the sequences. Our goal here is to allow tradeoff between the degeneracy and the coverag There are various reasons for studying DPD. Fo example, we were involved in a project with the groups of H.Lehrach (MPI Berlin) and D.Lancet (Weizmann) fo finding new human olfactory receptor (OR) genes. At the outset of the project (which preceded the publication of the human genome), only 127 OR genes were known, and the goal was to selectively amplify additional OR genes using degenerate primers. The rationale was that primers which match many of the known genes, would also amplify many new genes from the same family as well, whose sequences are closely related. Most OR genes contain conserved regions, and so the primer vould be designed to match such regions. OR genes are a A PCR (Polymerase Chain Reaction) primer sequence is single 1000bp exon, so amplification can be done on the called degenerate if some of its positions have sever nomic sequence. In gene families that contain introns possible bases (Kwok et al., 1994). For example, in the same technique can be applied to selectively amplify primer GG[CG]A[CTG]A the third position is C or G and cDNAs. The technique can be applied to various familie the fifth is C, T or G. The degeneracy of the primer is and to extracting genes from a particular family in an the number of unique sequence combinations it contains unsequenced species based on the known sequences o For example, the degeneracy of the above primer is 6. family members in a related species. In cDNA analysis Degenerate primers are as easy and cheap to produce one can use degenerate primers for amplifying and then as regular unique primers, are useful for amplifying suring frequencies of members of a gene family. several related genomic sequences, and have been used DPD is related to the Primer Selection Problem (Pearson in various applications. Most extant applications use low t al., 1996), in which the goal is to minimize the

degeneracies of up to hundreds. In this paper we study the problem of designing primers of high degeneracy. In a typical situation, one has a collection of related target sequences, , DNA sequences of homologous genes, and the goal is to design primers that will match as many of them as possible. A naïve solution would be to align large gene families, the number of primers needed to cover

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number of (non-degenerate) primers required to amplify

a set of DNA sequences. Several algorithms have been

developed to solve this problem, and some take into account various biological considerations and technical

constraints (see, e.g., Doi and Imai, 1997). However, for

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Nonoblivious 2-Opt heuristics for the traveling salesman problem. Approximation Limits of Linear Programs (Beyond Hierarchies). Validation of Distributed SDN Control Plane Under Uncertain Failures, IEEE/ACM Transactions on Networking, 27:3, (1234-1247), Online publication date: 1-Jun-2019.Brucato M, Abouzied A and Meliou A 2019. The book is organized around central algorithmic techniques for designing approximation algorithms, including greedy and local search algorithms, dynamic programming, linear and semidefinite programming, and randomization. Integrality gaps for strengthened linear relaxations of capacitated facility location, Mathematical Programming; Series A and B, 158:1-2, (99-141), Online publication date: 1-Jul-2016.D'Angelo G, Severini L and Velaj Y 2016. Thus unless P = NP, there are no efficient algorithms to find optimal solutions to such problems. 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NLopt includes implementations of a number of different optimization algorithms. These algorithms. These algorithms are listed below, including links to the original source code (if any) and citations to the relevant articles in the literature (see Citing NLopt)... Even where I found available free/open-source code for the various algorithms, I modified the code at least slightly (and in some cases ... Numerical analysis is the study of algorithms that use numerical analysis (as distinguished from discrete mathematics). Numerical analysis finds application in all fields of engineering and the physical sciences, and in the 21st century also the life and social sciences, medicine, business ... NPTEL provides E-learning through online Web and Video courses various streams. 01/09/2021 · State design pattern is used when an Object changes its behavior based on its internal state. If we have to change behavior of an object based on its state, we can have a state variable in the Object and use if-else condition block to perform different actions based on the state. Design and Analysis of Algorithms. Stanford University, Winter 2022. Instructors: Nima Anari and Moses Charikar Time: Mon & Wed 9:45 am - 11:15 am Location: Zoom for the first three weeks, then NVIDIA Auditorium Course Description: This course will cover the basic approaches and mindsets for analyzing and designing algorithms. Topics include the ... Course Description: This course introduces basic elements of the design and analysis of computer algorithms. Topics include the ... Course Description: This course introduces basic elements of the design and analysis of computer algorithms. Topics include the ... Course Description: This course introduces basic elements of the design and analysis of computer algorithms. Topics include the ... Course Description: This course will cover the basic approaches and mindsets for analyzing and conquer strategy, greedy methods, dynamic programming, basic graph algorithms, NP-completeness, and approximation algorithms. 21/11/2019 · Loop Optimization is the process of increasing execution speed and reducing the overheads associated with loops. It plays an important role in improving cache performance and making effective use of parallel processing capabilities. Most execution time of a scientific program is spent on loops. 07/06/2015 · Algorithms Based on Piecewise Linear Approximations to solve complicated problems for optimal solutions. Branch and Refine. The branch and refine algorithm is based on the piecewise linear approximation. It is an efficient way to solve a problem for the global ... Theory@CS.CMU. Carnegie Mellon University has a strong and diverse group in Algorithms and Complexity Theory. We try to provide a mathematical understanding of fundamental issues in Computer Science, and to use this understanding to produce better algorithms, protocols, and systems, as well as identify the inherent limitations of efficient computation. Course Description: This course introduces basic elements of the design and analysis, divide and conquer strategy, greedy methods, dynamic programming, basic graph algorithms, NP-completeness, and approximation algorithms. 29/07/2021 · Design an online book reader system (Object-Oriented Design). Asked In: Amazon, Microsoft, and many more interviews. Solution: Let's assume we want to design a basic online reading system that provides the following functionality: • Searching the database of books and reading a book. • User membership creation and extension

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