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An Integer Programming Approach to the Student-Project Allocation Problem with Preferences over Projects*

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Abstract. The Student-Project Allocation problem with preferences over Projects (SPA-P) involves sets of students, projects and lecturers, where the students and lecturers each have preferences over the projects. In this context, we typically seek a stable matching of students to projects (and lecturers). However, these stable matchings can have different sizes, and the problem of finding a maximum stable matching (MAX-SPA-P) is NP-hard. There are two known approximation algorithms for MAX-SPA-P, with performance guarantees of 2 and $\frac{3}{2}$. In this paper, we describe an Integer Programming (IP) model to enable MAX-SPA-P to be solved optimally. Following this, we present results arising from an empirical analysis that investigates how the solution produced by the approximation algorithms compares to the optimal solution obtained from the IP model, with respect to the size of the stable matchings constructed, on instances that are both randomly-generated and derived from real datasets. Our main finding is that the $\frac{3}{2}$ -approximation algorithm finds stable matchings that are very close to having maximum cardinality.

1 Introduction

Matching problems, which generally involve the assignment of a set of agents to another set of agents based on preferences, have wide applications in many real-world settings. One such application can be seen in an educational context, e.g., the allocation of pupils to schools, school-leavers to universities and students to projects. In the context of allocating students to projects, university lecturers propose a range of projects, and each student is required to provide a preference over the available projects that she finds acceptable. Lecturers may also have preferences over the students that find their project acceptable and/or the projects that they offer. There may also be upper bounds on the number of

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students that can be assigned to a particular project, and the number of students that a given lecturer is willing to supervise. The problem then is to allocate students to projects based on these preferences and capacity constraints – the so-called *Student-Project Allocation problem* (SPA) [3,11].

Two major models of SPA exist in the literature: one permits preferences only from the students [2,6,10,14], while the other permits preferences from the students and lecturers [1,8]. Given the large number of students that are typically involved in such an allocation process, many university departments seek to automate the allocation of students to projects. Examples include the School of Computing Science, University of Glasgow [10], the Faculty of Science, University of Southern Denmark [4], the Department of Computing Science, University of York [8], and elsewhere [2,3,6,16].

In general, we seek a *matching*, which is a set of agent pairs who find one another acceptable that satisfies the capacities of the agents involved. For matching problems where preferences exist from the two sets of agents involved (e.g., junior doctors and hospitals in the classical *Hospitals-Residents problem* (HR) [5], or students and lecturers in the context of SPA), it has been argued that the desired property for a matching one should seek is that of *stability* [15]. Informally, a *stable matching* ensures that no acceptable pair of agents who are not matched together would rather be assigned to each other than remain with their current assignees.

Abraham, Irving and Manlove [1] proposed two linear-time algorithms to find a stable matching in a variant of SPA where students have preferences over projects, whilst lecturers have preferences over students. The stable matching produced by the first algorithm is student-optimal (that is, students have the best possible projects that they could obtain in any stable matching) while the one produced by the second algorithm is lecturer-optimal (that is, lecturers have the best possible students that they could obtain in any stable matching).

Manlove and O'Malley [12] proposed another variant of SPA where both students and lecturers have preferences over projects, referred to as SPA-P. In their paper, they formulated an appropriate stability definition for SPA-P, and they showed that stable matchings in this context can have different sizes. Moreover, in addition to stability, a very important requirement in practice is to match as many students to projects as possible. Consequently, Manlove and O'Malley [12] proved that the problem of finding a maximum cardinality stable matching, denoted MAX-SPA-P, is NP-hard. Further, they gave a polynomial-time 2-approximation algorithm for MAX-SPA-P. Subsequently, Iwama, Miyazaki and Yanagisawa [7] described an improved approximation algorithm with an upper bound of $\frac{3}{2}$, which builds on the one described in [12]. In addition, Iwama et al. [7] showed that MAX-SPA-P is not approximable within $\frac{21}{19} - \epsilon$, for any $\epsilon > 0$, unless P = NP. For the upper bound, they modified Manlove and O'Malley's algorithm [12] using Király's idea [9] for the approximation algorithm to find a maximum stable matching in a variant of the Stable Marriage problem.

Considering the fact that the existing algorithms for MAX-SPA-P are only guaranteed to produce an approximate solution, we seek another technique to

enable MAX-SPA-P to be solved optimally. Integer Programming (IP) is a powerful technique for producing optimal solutions to a range of NP-hard optimisation problems, with the aid of commercial optimisation solvers, e.g., Gurobi [17], GLPK [18] and CPLEX [19]. These solvers can allow IP models to be solved in a reasonable amount of time, even with respect to problem instances that occur in practical applications.

Our Contribution. In Sect. 3, we describe an IP model to enable MAX-SPA-P to be solved optimally, and present a correctness result. In Sect. 4, we present results arising from an empirical analysis that investigates how the solution produced by the approximation algorithms compares to the optimal solution obtained from our IP model, with respect to the size of the stable matchings constructed, on instances that are both randomly-generated and derived from real datasets. These real datasets are based on actual student preference data and manufactured lecturer preference data from previous runs of student-project allocation processes at the School of Computing Science, University of Glasgow. We also present results showing the time taken by the IP model to solve the problem instances optimally. Our main finding is that the $\frac{3}{2}$ -approximation algorithm finds stable matchings that are very close to having maximum cardinality. The next section gives a formal definition for SPA-P.

2 Definitions and Preliminaries

We give a formal definition for SPA-P as described in the literature [12]. An instance I of SPA-P involves a set $\mathcal{S} = \{s_1, s_2, \ldots, s_{n_1}\}$ of students, a set $\mathcal{P} = \{p_1, p_2, \ldots, p_{n_2}\}$ of projects and a set $\mathcal{L} = \{l_1, l_2, \ldots, l_{n_3}\}$ of lecturers. Each lecturer $l_k \in \mathcal{L}$ offers a non-empty subset of projects, denoted by P_k . We assume that $P_1, P_2, \ldots, P_{n_3}$ partitions \mathcal{P} (that is, each project is offered by one lecturer). Also, each student $s_i \in \mathcal{S}$ has an acceptable set of projects $A_i \subseteq \mathcal{P}$. We call a pair $(s_i, p_j) \in \mathcal{S} \times \mathcal{P}$ an acceptable pair if $p_j \in A_i$. Moreover s_i ranks A_i in strict order of preference. Similarly, each lecturer l_k ranks P_k in strict order of preference. Finally, each project $p_j \in \mathcal{P}$ and lecturer $l_k \in \mathcal{L}$ has a positive capacity denoted by c_j and d_k respectively.

An assignment M is a subset of $S \times P$ where $(s_i, p_j) \in M$ implies that s_i finds p_j acceptable (that is, $p_j \in A_i$). We define the size of M as the number of (student, project) pairs in M, denoted |M|. If $(s_i, p_j) \in M$, we say that s_i is assigned to p_j and p_j is assigned s_i . Furthermore, we denote the project assigned to student s_i in M as $M(s_i)$ (if s_i is unassigned in M then $M(s_i)$ is undefined). Similarly, we denote the set of students assigned to project p_j in M as $M(p_j)$. For ease of exposition, if s_i is assigned to a project p_j offered by lecturer l_k , we may also say that s_i is assigned to l_k , and l_k is assigned s_i . Thus we denote the set of students assigned to l_k in M as $M(l_k)$.

A project $p_j \in \mathcal{P}$ is full, undersubscribed or oversubscribed in M if $|M(p_j)|$ is equal to, less than or greater than c_j , respectively. The corresponding terms apply to each lecturer l_k with respect to d_k . We say that a project $p_j \in \mathcal{P}$ is non-empty if $|M(p_j)| > 0$.

A matching M is an assignment such that $|M(s_i)| \leq 1$ for each $s_i \in \mathcal{S}$, $|M(p_j)| \leq c_j$ for each $p_j \in \mathcal{P}$, and $|M(l_k)| \leq d_k$ for each $l_k \in \mathcal{L}$ (that is, each student is assigned to at most one project, and no project or lecturer is oversubscribed). Given a matching M, an acceptable pair $(s_i, p_j) \in (\mathcal{S} \times \mathcal{P}) \setminus M$ is a blocking pair of M if the following conditions are satisfied:

- 1. either s_i is unassigned in M or s_i prefers p_j to $M(s_i)$, and p_j is undersubscribed, and either
 - (a) $s_i \in M(l_k)$ and l_k prefers p_j to $M(s_i)$, or
 - (b) $s_i \notin M(l_k)$ and l_k is undersubscribed, or
 - (c) $s_i \notin M(l_k)$ and l_k prefers p_j to his worst non-empty project, where l_k is the lecturer who offers p_j .

If such a pair were to occur, it would undermine the integrity of the matching as the student and lecturer involved would rather be assigned together than remain in their current assignment. With respect to the spa-P instance given in Fig. 1, $M_1 = \{(s_1, p_3), (s_2, p_1)\}$ is clearly a matching. It is obvious that each of students s_1 and s_2 is matched to her first ranked project in M_1 . Although s_3 is unassigned in M_1 , the lecturer offering p_3 (the only project that s_3 finds acceptable) is assumed to be indifferent among those students who find p_3 acceptable. Also p_3 is full in M_1 . Thus, we say that M_1 admits no blocking pair.

Fig. 1. An instance I_1 of SPA-P. Each project has capacity 1, whilst each of lecturer l_1 and l_2 has capacity 2 and 1 respectively.

Another way in which a matching could be undermined is through a group of students acting together. Given a matching M, a coalition is a set of students $\{s_{i_0},\ldots,s_{i_{r-1}}\}$, for some $r\geq 2$ such that each student s_{i_j} $(0\leq j\leq r-1)$ is assigned in M and prefers $M(s_{i_{j+1}})$ to $M(s_{i_j})$, where addition is performed modulo r. With respect to Fig. 1, the matching $M_2=\{(s_1,p_1),(s_2,p_2),(s_3,p_3)\}$ admits a coalition $\{s_1,s_2\}$, as students s_1 and s_2 would rather permute their assigned projects in M_2 so as to be better off. We note that the number of students assigned to each project and lecturer involved in any such swap remains the same after such a permutation. Moreover, the lecturers involved would have no incentive to prevent the switch from occurring since they are assumed to be indifferent between the students assigned to the projects they are offering. If a matching admits no coalition, we define such matching to be coalition-free.

Given an instance I of SPA-P, we define a matching M in I to be *stable* if M admits no blocking pair and is coalition-free. It turns out that with respect to this definition, stable matchings in I can have different sizes. Clearly, each

of the matchings $M_1 = \{(s_1, p_3), (s_2, p_1)\}\$ and $M_3 = \{(s_1, p_2), (s_2, p_1), (s_3, p_3)\}\$ is stable in the SPA-P instance I₁ shown in Fig. 1. The varying sizes of the stable matchings produced naturally leads to the problem of finding a maximum cardinality stable matching given an instance of SPA-P, which we denote by MAX-SPA-P. In the next section, we describe our IP model to enable MAX-SPA-P to be solved optimally.

An IP model for MAX-SPA-P

Let I be an instance of SPA-P involving a set $S = \{s_1, s_2, \dots, s_{n_1}\}$ of students, a set $\mathcal{P} = \{p_1, p_2, \dots, p_{n_2}\}$ of projects and a set $\mathcal{L} = \{l_1, l_2, \dots, l_{n_3}\}$ of lecturers. We construct an IP model J of I as follows. Firstly, we create binary variables $x_{i,j} \in \{0,1\}$ $(1 \le i \le n_1, 1 \le j \le n_2)$ for each acceptable pair $(s_i, p_j) \in$ $\mathcal{S} \times \mathcal{P}$ such that $x_{i,j}$ indicates whether s_i is assigned to p_j in a solution or not. Henceforth, we denote by S a solution in the IP model J, and we denote by Mthe matching derived from S. If $x_{i,j} = 1$ under S then intuitively s_i is assigned to p_i in M, otherwise s_i is not assigned to p_i in M. In what follows, we give the constraints to ensure that the assignment obtained from a feasible solution in J is a matching.

Matching Constraints. The feasibility of a matching can be ensured with the following three sets of constraints.

$$\sum_{p_j \in A_i} x_{i,j} \le 1 \qquad (1 \le i \le n_1), \tag{1}$$

$$\sum_{i=1}^{n_1} x_{i,j} \le c_j \qquad (1 \le j \le n_2), \tag{2}$$

$$\sum_{i=1}^{n_1} x_{i,j} \le c_j \qquad (1 \le j \le n_2),$$

$$\sum_{i=1}^{n_1} \sum_{p_j \in P_k} x_{i,j} \le d_k \qquad (1 \le k \le n_3) .$$
(3)

Note that (1) implies that each student $s_i \in \mathcal{S}$ is not assigned to more than one project, while (2) and (3) implies that the capacity of each project $p_i \in \mathcal{P}$ and each lecturer $l_k \in \mathcal{L}$ is not exceeded.

We define $rank(s_i, p_i)$, the rank of p_i on s_i 's preference list, to be r+1 where r is the number of projects that s_i prefers to p_i . An analogous definition holds for $rank(l_k, p_j)$, the rank of p_j on l_k 's preference list. With respect to an acceptable pair (s_i, p_j) , we define $S_{i,j} = \{p_{j'} \in A_i : rank(s_i, p_{j'}) \leq rank(s_i, p_j)\}$, the set of projects that s_i likes as much as p_j . For a project p_j offered by lecturer $l_k \in \mathcal{L}$, we also define $T_{k,j} = \{p_q \in P_k : rank(l_k, p_j) < rank(l_k, p_q)\}$, the set of projects that are worse than p_i on l_k 's preference list.

In what follows, we fix an arbitrary acceptable pair (s_i, p_j) and we impose constraints to ensure that (s_i, p_i) is not a blocking pair of the matching M (that is, (s_i, p_j) is not a type 1(a), type 1(b) or type 1(c) blocking pair of M). Firstly, let l_k be the lecturer who offers p_j .

Blocking Pair Constraints. We define $\theta_{i,j} = 1 - \sum_{p_{j'} \in S_{i,j}} x_{i,j'}$. Intuitively, $\theta_{i,j} = 1$ if and only if s_i is unassigned in M or prefers p_j to $M(s_i)$. Next we create a binary variable α_j in J such that $\alpha_j = 1$ corresponds to the case when p_j is undersubscribed in M. We enforce this condition by imposing the following constraint.

$$c_j \alpha_j \ge c_j - \sum_{i'=1}^{n_1} x_{i',j}$$
, (4)

where $\sum_{i'=1}^{n_1} x_{i',j} = |M(p_j)|$. If p_j is undersubscribed in M then the RHS of (4) is at least 1, and this implies that $\alpha_j = 1$. Otherwise, α_j is not constrained. Now let $\gamma_{i,j,k} = \sum_{p_{j'} \in T_{k,j}} x_{i,j'}$. Intuitively, if $\gamma_{i,j,k} = 1$ in S then s_i is assigned to a project $p_{j'}$ offered by l_k in M, where l_k prefers p_j to $p_{j'}$. The following constraint ensures that (s_i, p_j) does not form a type 1(a) blocking pair of M.

$$\theta_{i,j} + \alpha_j + \gamma_{i,j,k} \le 2 . \tag{5}$$

Note that if the sum of the binary variables in the LHS of (5) is less than or equal to 2, this implies that at least one of the variables, say $\gamma_{i,j,k}$, is 0. Thus the pair (s_i, p_j) is not a type 1(a) blocking pair of M.

Next we define $\beta_{i,k} = \sum_{p_{j'} \in P_k} x_{i,j'}$. Clearly, s_i is assigned to a project offered by l_k in M if and only if $\beta_{i,k} = 1$ in S. Now we create a binary variable δ_k in J such that $\delta_k = 1$ in S corresponds to the case when l_k is undersubscribed in M. We enforce this condition by imposing the following constraint.

$$d_k \delta_k \ge d_k - \sum_{i'=1}^{n_1} \sum_{p_{i'} \in P_k} x_{i',j'} ,$$
 (6)

where $\sum_{i'=1}^{n_1} \sum_{p_{j'} \in P_k} x_{i',j'} = |M(l_k)|$. If l_k is undersubscribed in M then the RHS of (6) is at least 1, and this implies that $\delta_k = 1$. Otherwise, δ_k is not constrained. The following constraint ensures that (s_i, p_j) does not form a type 1(b) blocking pair of M.

$$\theta_{i,j} + \alpha_j + (1 - \beta_{i,k}) + \delta_k \le 3 .$$
(7)

We define $D_{k,j} = \{p_{j'} \in P_k : rank(l_k, p_{j'}) \leq rank(l_k, p_j)\}$, the set of projects that l_k likes as much as p_j . Next, we create a binary variable $\eta_{j,k}$ in J such that $\eta_{j,k} = 1$ if l_k is full and prefers p_j to his worst non-empty project in S. We enforce this by imposing the following constraint.

$$d_k \eta_{j,k} \ge d_k - \sum_{i'=1}^{n_1} \sum_{p_{j'} \in D_{k,j}} x_{i',j'} . \tag{8}$$

Finally, to avoid a type 1(c) blocking pair, we impose the following constraint.

$$\theta_{i,j} + \alpha_j + (1 - \beta_{i,k}) + \eta_{j,k} \le 3 .$$

$$(9)$$

Next, we give the constraints to ensure that the matching obtained from a feasible solution in J is coalition-free.

Coalition Constraints. First, we introduce some additional notation. Given an instance I' of SPA-P and a matching M' in I', we define the envy graph G(M') = (S, A), where the vertex set S is the set of students in I', and the arc set $A = \{(s_i, s_{i'}) : s_i \text{ prefers } M'(s_{i'}) \text{ to } M'(s_i)\}$. It is clear that the matching $M_2 = \{(s_1, p_1), (s_2, p_2), (s_3, p_3)\}$ admits a coalition $\{s_1, s_2\}$ with respect to the instance given in Fig. 1. The resulting envy graph $G(M_2)$ is illustrated below.

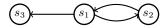


Fig. 2. The envy graph $G(M_2)$ corresponding to the SPA-P instance in Fig. 1.

Clearly, G(M') contains a directed cycle if and only if M' admits a coalition. Moreover, G(M') is acyclic if and only if it admits a topological ordering. Now to ensure that the matching M obtained from a feasible solution S under J is coalition-free, we will enforce J to encode the envy graph G(M) and impose the condition that it must admit a topological ordering. In what follows, we build on our IP model J of I.

We create a binary variable $e_{i,i'}$ for each $(s_i, s_{i'}) \in \mathcal{S} \times \mathcal{S}$, $s_i \neq s_{i'}$, such that the $e_{i,i'}$ variables will correspond to the adjacency matrix of G(M). For each i and i' $(1 \leq i \leq n_1, 1 \leq i' \leq n_1, i \neq i')$ and for each j and j' $(1 \leq j \leq n_2, 1 \leq j' \leq n_2)$ such that s_i prefers $p_{j'}$ to p_j , we impose the following constraint.

$$e_{i,i'} + 1 \ge x_{i,j} + x_{i',j'} \tag{10}$$

If $(s_i, p_j) \in M$ and $(s_{i'}, p_{j'}) \in M$ and s_i prefers $p_{j'}$ to p_j , then $e_{i,i'} = 1$ and we say s_i envies $s_{i'}$. Otherwise, $e_{i,i'}$ is not constrained. Next we enforce the condition that G(M) must have a topological ordering. To hold the label of each vertex in a topological ordering, we create an integer-valued variable v_i corresponding to each student $s_i \in \mathcal{S}$ (and intuitively to each vertex in G(M)). We wish to enforce the constraint that if $e_{i,i'} = 1$ (that is, $(s_i, s_{i'}) \in A$), then $v_i < v_{i'}$ (that is, the label of vertex s_i is smaller than the label of vertex $s_{i'}$). This is achieved by imposing the following constraint for all i and i' $(1 \le i \le n_1, 1 \le i' \le n_1, i \ne i')$.

$$v_i < v_{i'} + n_1(1 - e_{i,i'})$$
 (11)

Note that the LHS of (11) is strictly less than the RHS of (11) if and only if G(M) does not admit a directed cycle, and this implies that M is coalition-free.

Variables. We define a collective notation for each variable involved in J as follows.

$$\begin{split} X &= \{x_{i,j}: 1 \leq i \leq n_1, 1 \leq j \leq n_2\}, \quad \Lambda = \{\alpha_j: 1 \leq j \leq n_2\}, \\ H &= \{\eta_{j,k}: 1 \leq j \leq n_2, 1 \leq k \leq n_3\}, \quad \Delta = \{\delta_k: 1 \leq k \leq n_3\}, \\ E &= \{e_{i,i'}: 1 \leq i \leq n_1, 1 \leq i' \leq n_1\}, \quad V = \{v_i: 1 \leq i \leq n_1\} \ . \end{split}$$

Objective Function. The objective function given below is a summation of all the $x_{i,j}$ binary variables. It seeks to maximize the number of students assigned (that is, the cardinality of the matching).

$$\max \sum_{i=1}^{n_1} \sum_{p_j \in A_i} x_{i,j} .$$
 (12)

Finally, we have constructed an IP model J of I comprising the set of integer-valued variables X, Λ, H, Δ, E and V, the set of constraints (1) - (11) and an objective function (12). Note that J can then be used to solve MAX-SPA-P optimally. Given an instance I of SPA-P formulated as an IP model J using the above transformation, we present the following result regarding the correctness of J (see [13] for proof).

Theorem 1. A feasible solution to J is optimal if and only if the corresponding stable matching in I is of maximum cardinality.

4 Empirical Analysis

In this section we present results from an empirical analysis that investigates how the sizes of the stable matchings produced by the approximation algorithms compares to the optimal solution obtained from our IP model, on SPA-P instances that are both randomly-generated and derived from real datasets.

4.1 Experimental Setup

There are clearly several parameters that can be varied, such as the number of students, projects and lecturers; the length of the students' preference lists; as well as the total capacities of the projects and lecturers. For each range of values for the first two parameters, we generated a set of random SPA-P instances. In each set, we record the average size of a stable matching obtained from running the approximation algorithms and the IP model. Further, we consider the average time taken for the IP model to find an optimal solution.

By design, the approximation algorithms were randomised with respect to the sequence in which students apply to projects, and the choice of students to reject when projects and/or lecturers become full. In the light of this, for each dataset, we also run the approximation algorithms 100 times and record the size of the largest stable matching obtained over these runs. Our experiments therefore involve five algorithms: the optimal IP-based algorithm, the two approximation algorithms run once, and the two approximation algorithms run 100 times.

We performed our experiments on a machine with dual Intel Xeon CPU E5-2640 processors with 64GB of RAM, running Ubuntu 14.04. Each of the approximation algorithms was implemented in Java¹. For our IP model, we carried

¹ https://github.com/sofiat-olaosebikan/spa-p-isco-2018

out the implementation using the Gurobi optimisation solver in Java¹. For correctness testing on these implementations, we designed a stability checker which verifies that the matching returned by the approximation algorithms and the IP model does not admit a blocking pair or a coalition.

4.2 Experimental Results

Randomly-generated Datasets. All the SPA-P instances we randomly generated involved n_1 students (n_1 is henceforth referred to as the size of the instance), $0.5n_1$ projects, $0.2n_1$ lecturers and $1.1n_1$ total project capacity which was randomly distributed amongst the projects. The capacity for each lecturer l_k was chosen randomly to lie between the highest capacity of the projects offered by l_k and the sum of the capacities of the projects that l_k offers. In the first experiment, we present results obtained from comparing the performance of the IP model, with and without the coalition constraints in place.

Experiment 0. We increased the number of students n_1 while maintaining a ratio of projects, lecturers, project capacities and lecturer capacities as described above. For various values of n_1 ($100 \le n_1 \le 1000$) in increments of 100, we created 100 randomly-generated instances. Each student's preference list contained a minimum of 2 and a maximum of 5 projects. With respect to each value of n_1 , we obtained the average time taken for the IP solver to output a solution, both with and without the coalition constraints being enforced. The results, displayed in Table 1, show that when we removed the coalition constraints, the average time for the IP solver to output a solution is significantly faster than when we enforced the coalition constraints.

In the remaining experiments, we thus remove the constraints that enforce the absence of a coalition in the solution. We are able to do this for the purposes of these experiments because the largest size of a stable matching is equal to the largest size of a matching that potentially admits a coalition but admits no blocking pair², and we were primarily concerned with measuring stable matching cardinalities. However the absence of the coalition constraints should be borne in mind when interpreting the IP solver runtime data in what follows.

In the next two experiments, we discuss results obtained from running the five algorithms on randomly-generated datasets.

Experiment 1. As in the previous experiment, we maintained the ratio of the number of students to projects, lecturers and total project capacity; as well as the length of the students' preference lists. For various values of n_1 ($100 \le n_1 \le 2500$) in increments of 100, we created 1000 randomly-generated instances. With respect to each value of n_1 , we obtained the average sizes of stable matchings constructed by the five algorithms run over the 1000 instances. The result displayed in Fig. 3 (and also in Fig. 4) shows the ratio of the average size of the

 $^{^2}$ This holds because the number of students assigned to each project and lecturer in the matching remains the same even after the students involved in such coalition permute their assigned projects.

stable matching produced by the approximation algorithms with respect to the maximum cardinality matching produced by the IP solver.

Figure 3 shows that each of the approximation algorithms produces stable matchings with a much higher cardinality from multiple runs, compared to running them only once. Also, the average time taken for the IP solver to find a maximum cardinality matching increases as the size of the instance increases, with a running time of less than one second for instance size 100, increasing roughly linearly to 13 seconds for instance size 2500 (see [13, Fig. 3(b)]).

Experiment 2. In this experiment, we varied the length of each student's preference list while maintaining a fixed number of students, projects, lecturers and total project capacity. For various values of x ($2 \le x \le 10$), we generated 1000 instances, each involving 1000 students, with each student's preference list containing exactly x projects. The result for all values of x is displayed in Fig. 4. Figure 4 shows that as we increase the preference list length, the stable matchings produced by each of the approximation algorithms gets close to having maximum cardinality. It also shows that with a preference list length greater than 5, the $\frac{3}{2}$ -approximation algorithm produces an optimal solution, even on a single run. Moreover, the average time taken for the IP solver to find a maximum matching increases as the length of the students' preference lists increases, with a running time of two seconds when each student's preference list is of length 2, increasing roughly linearly to 17 seconds when each student's preference list is of length 10 (see [13, Fig. 4(b)]).

Real Datasets. The real datasets in this paper are based on actual student preference data and manufactured lecturer data from previous runs of student-project allocation processes at the School of Computing Science, University of Glasgow. Table 2 shows the properties of the real datasets, where n_1, n_2 and n_3 denotes the number of students, projects and lecturers respectively; and l denotes the length of each student's preference list. For all these datasets, each project has a capacity of 1. In the next experiment, we discuss how the lecturer preferences were generated. We also discuss the results obtained from running the five algorithms on the corresponding SPA-P instances.

Experiment 3. We derived the lecturer preference data from the real datasets as follows. For each lecturer l_k , and for each project p_j offered by l_k , we obtained the number a_j of students that find p_j acceptable. Next, we generated a strict preference list for l_k by arranging l_k 's proposed projects in (i) a random manner, (ii) ascending order of a_j , and (iii) descending order of a_j , where (ii) and (iii) are taken over all projects that l_k offers. Table 2 shows the size of stable matchings obtained from the five algorithms, where A, B, C, D and E denotes the solution obtained from the IP model, 100 runs of $\frac{3}{2}$ -approximation algorithm, single run of $\frac{3}{2}$ -approximation algorithm respectively. The results are essentially consistent with the findings in the previous experiments, that is, the $\frac{3}{2}$ -approximation algorithm produces stable matchings whose sizes are close to optimal.

Table 1. Results for Experiment 0. Average time (in seconds) for the IP solver to output a solution, both with and without the coalition constraints being enforced.

Size of instance	100	200	300	400	500	600	700	800	900	1000
Av. time without coalition	0.12	0.27	0.46	0.69	0.89	1.17	1.50	1.86	2.20	2.61
Av. time with coalition	0.71	2.43	4.84	9.15	13.15	19.34	28.36	38.18	48.48	63.50

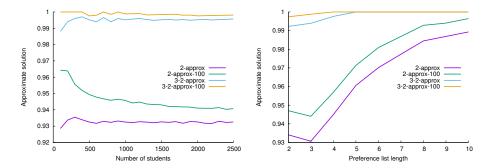


Fig. 3. Result for Experiment 1.

Fig. 4. Result for Experiment 2.

Table 2. Properties of the real datasets and results for Experiment 3.

					Random					Most popular					Least popular				
Year	n_1	n_2	n_3	l	A	B	C	D	E	A	B	C	D	E	A	B	C	D	E
2014	55	149	38	6	55	55	55	54	53	55	55	55	54	50	55	55	55	54	52
2015	76	197	46	6	76	76	76	76	72	76	76	76	76	72	76	76	76	76	75
2016	92	214	44	6	84	82	83	77	75	85	85	83	79	76	82	80	77	76	74
2017	90	289	59	4	89	87	85	80	76	90	89	86	81	79	88	85	84	80	77

4.3 Discussions and Concluding Remarks

The results presented in this section suggest that even as we increase the number of students, projects, lecturers, and the length of the students' preference lists, each of the approximation algorithms finds stable matchings that are close to having maximum cardinality, outperforming their approximation factor. Perhaps most interesting is the $\frac{3}{2}$ -approximation algorithm, which finds stable matchings that are very close in size to optimal, even on a single run. These results also holds analogously for the instances derived from real datasets.

We remark that when we removed the coalition constraints, we were able to run the IP model on an instance size of 10000, with the solver returning a maximum matching in an average time of 100 seconds, over 100 randomly-generated instances. This shows that the IP model (without enforcing the coalition constraints), can be run on SPA-P instances that appear in practice, to find maximum cardinality matchings that admit no blocking pair. Coalitions should then

be eliminated in polynomial time by repeatedly constructing an *envy graph*, similar to the one described in [11, p.290], finding a directed cycle and letting the students in the cycle swap projects.

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