

# Matching with Couples: Stability and Incentives in Large Markets

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## Abstract

Complementarities pose problems in models of two-sided matching markets. This has been a longstanding issue in the design of centralized labor market clearinghouses that need to accommodate complementarities due to couples, as in the US market for medical doctors. These clearinghouses aim for a stable matching but a stable matching does not necessarily exist when couples are present. This paper provides conditions under which a stable matching exists with high probability in large markets. Moreover, we present a mechanism that finds a stable matching with high probability and in which truth-telling by all participants is an approximate equilibrium. We relate these theoretical results to data from the labor market for clinical psychologists, in which stable matchings exist for all years of our data, despite the presence of couples.

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# 1 Introduction

One of the big 20th century transformations of the American labor market involves the increased labor force participation of married women, and the consequent growth in the number of two-career households.<sup>1</sup> When a couple needs two jobs, they face a hard problem of coordination with each other and with their prospective employers. The search and matching process for the spouses can involve very different timing of searches and hiring. The couple may be forced to make a decision on a job offer for one member of the couple before knowing what complementary jobs may become available for the other or what better pairs of jobs might become available elsewhere.

An unusually clear view of this problem can be found in the history of the entry-level labor market for American doctors. Since the early 1900s, new U.S. medical graduates have been first employed as “residents” at hospitals, where they work under the supervision of more senior, licensed doctors. This market experienced serious problems having to do with the timing of offers and acceptances, and this *unraveling* of the market led to the creation of a centralized clearinghouse in the 1950s that drew high rates of participation (see Roth (1984) and Roth and Xing (1994) for further details). Medical graduates were almost all men throughout this period, but by the 1970s there were enough women graduating from medical school so that it was not unheard of for two new medical graduates to be married to each other. Many couples felt that the existing clearinghouse did not serve them well, and starting in the 1970s, significant numbers of these couples began seeking jobs outside of the clearinghouse.

Roth (1984) argues that this was because the existing matching algorithm did not allow couples to appropriately express preferences. That paper shows that, in a market without couples, the 1950s clearinghouse algorithm is equivalent to the deferred acceptance algorithm of Gale and Shapley (1962), and that it produces a *stable* matching – loosely speaking, this is a matching such that there is no pair of hospital and doctor who want to be matched with each other rather than accepting the prescribed matching.<sup>2</sup> It then observes that the algorithm often fails to find a stable matching when there are couples, and argues that a main problem of the mechanism is that it does not allow couples to report preferences over *pairs* of positions, one for each member of the couple. Roth and Peranson (1999) offer an algorithm that elicits couples’ preferences over pairs of positions, and this design has been adopted by more than 40 centralized clearinghouses such as the medical market for American doctors, the National Resident Matching Program (NRMP).<sup>3</sup>

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<sup>1</sup>See, for instance, Costa and Kahn (2000) for a description of the trends in the labor market choices for college-educated couples since World War II.

<sup>2</sup>Section 3 provides a precise definition of our stability concept. The evidence suggests that the stability of the match plays an important role in attracting high rates of participation (Roth, 1990, 1991).

<sup>3</sup>See Roth (2008) for a recent list of these clearinghouses as well as a survey of the literature. See also Sönmez and

But the problem is difficult even if couples are allowed to express their preferences over pairs of positions, because there does not necessarily exist a stable matching in markets with couples (Roth, 1984). However, some matching clearinghouses seem to regularly entertain high rates of participation and appear to have produced matchings that are honored by participants. In fact, it has been reported that there have only been a few occasions in which a stable matching was not found over the last decade in several dozen annual markets (Peranson, private communication). Moreover, in the largest of these markets, the NRMP, Roth and Peranson (1999) run a number of matching algorithms using submitted preferences from 1993, 1994 and 1995 and find no instance in which any of these algorithms failed to produce a stable matching. Why do these matching clearinghouses produce stable outcomes even though existing theory suggests that stable matchings may not exist when couples are present?

This is the puzzle we address, and this paper argues that the answer may have to do with the size of the market. We consider a sequence of markets indexed by the number of hospitals, where doctors' preferences are drawn from some distribution. When the number of couples does not grow too fast relative to market size, under some regularity conditions, we demonstrate that the probability that a stable matching exists converges to one as the market size approaches infinity. Moreover, we provide an algorithm that finds a stable matching with a probability that converges to one as the market size approaches infinity.

In practice, preferences are private information, and the matching clearinghouse needs to elicit the information from participants. This motivates our analysis of the incentive properties of a particular matching mechanism in markets with couples. More specifically, we first define a mechanism similar to Roth and Peranson's algorithm, used in many existing markets. For a Bayesian game in which doctors and hospitals submit their preferences to this mechanism, we establish that truth-telling is an approximate Bayes Nash equilibrium in any market with a sufficiently large number of hospitals.

As our theoretical analysis only provides limit results, we study data on submitted preferences from the centralized market for clinical psychologists. In the late 1990s, the market evolved from a decentralized one to one employing a centralized clearinghouse (Roth and Xing, 1997), where a key design issue was whether it would be possible to accommodate the presence of couples. Keilin (1998) reports that under the old decentralized system couples had difficulties coordinating their internship choices. In 1999, clinical psychologists adopted a centralized clearinghouse using an algorithm based on Roth and Peranson (1999), where couples are allowed to express preferences over hospital pairs. We explore a variation of the Roth-Peranson procedure to investigate the existence of a stable matching with respect to the stated preferences of participants from nine years of data from 1999-2007. Using our algorithm, we are able to find a stable matching in all nine years.

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Ünver (2009).

## Related Literature

This paper is related to several lines of work. First, it is part of research in two-sided matching with couples. Existing studies are mostly negative: Roth (1984) and unpublished work by Sotomayor show that there does not necessarily exist a stable matching when there are couples. Klaus and Klijn (2005) provide a maximal domain of couple preferences that guarantees the existence of stable matchings. While their preference domain has a natural interpretation, our paper finds that preferences of almost all couples in our psychology market data violate their condition.<sup>4</sup> The present paper takes a new and different approach based on large market arguments to establish new positive results.

The second line of studies related to this paper is the growing literature on large matching markets. This paper uses large market arguments similar to Kojima and Pathak (2009). That paper as well as the current one are motivated by the analysis of Roth and Peranson (1999). They conduct simulations based on the NRMP data and randomly generated data and find suggestive evidence that a stable matching is likely to exist and stable matching mechanisms are difficult to manipulate in large markets. One of their findings is that the number of stable matchings becomes relatively small in large markets, which might suggest that if a stable matching does not exist in a finite market, then it is unlikely to exist in a large market. Our analysis finds, on the contrary, that largeness of the market tends to overcome non-existence in the finite market.

Recently, large market arguments are used by increasingly many studies to analyze incentive and efficiency of matching mechanisms, not only in two-sided matching (Bulow and Levin, 2006; Immorlica and Mahdian, 2005) but also in the closely related assignment or one-sided matching model (Abdulkadiroğlu et al., 2008; Che and Kojima, 2010; Kojima and Manea, 2009; Manea, 2009). However, the use of large market arguments to establish the existence of a stable matching is not found in these papers, and appears novel in the matching literature.

While the analysis of large markets is relatively new in the matching literature, it has a long tradition in economics. For example, Roberts and Postlewaite (1976) and Jackson and Manelli (1997) show that, under some conditions, the Walrasian mechanism is difficult to manipulate in large exchange economies. Similarly, Gresik and Satterthwaite (1989) and Rustichini et al. (1994) study incentive properties of a large class of double auction mechanisms.

Finally, a couple preference is a particular form of complementarity, and this paper can be put

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<sup>4</sup>Preference restrictions are also used to study incentives for manipulation in matching markets. The restriction under which incentive compatibility can be established is often very strong, as shown by Alcalde and Barberà (1994), Kesten (2006b), Kojima (2007), and Konishi and Ünver (2006b) for various kinds of manipulations. Similarly, in the context of resource allocation and school choice (Abdulkadiroğlu and Sönmez, 2003), necessary and sufficient conditions for desirable properties such as efficiency and incentive compatibility are strict (Ergin, 2002; Kesten, 2006a; Haeringer and Klijn, 2009; Ehlers and Erdil, 2009). Another approach is based on incomplete information (Roth and Rothblum, 1999; Ehlers, 2004, 2008; Kesten, 2009; Erdil and Ergin, 2008).

in the context of the larger research program on the role of complementarities in resource allocation. Complementarities have been identified to cause non-existence of desirable solutions in various contexts of resource allocation. There has been a recent flurry of investigations on complementarities and existence problems in auction markets (Milgrom, 2004; Gul and Stacchetti, 2000; Sun and Yang, 2009), general equilibrium with indivisible goods (Bikhchandani and Ostroy, 2002; Gul and Stacchetti, 1999; Sun and Yang, 2006), and matching markets (Hatfield and Kominers, 2009; Ostrovsky, 2008; Sönmez and Ünver, 2010; Pycia, 2010).

The layout of this paper is as follows. The next section describes some features of the market for clinical psychologists and lays out a series of stylized facts on matching with couples based on data from this market. Section 3 defines the model and describes a simple theory of matching with couples in a finite market. Section 4 introduces the large market assumptions. Section 5 states our main results on existence, while Section 6 describes our incentive result. Section 7 concludes.

## 2 The Market for Clinical Psychologists

### 2.1 Background

The story of how design has influenced and reacted to the presence of couples in the NRMP has parallels in the evolution of the market for clinical psychologists. Roth and Xing (1997) described the format of this market until the early 1990s. In the latter half of that decade, this market continued to operate in a decentralized fashion based on a “uniform notification day” system in which offers were given to internship applicants over the telephone within a specific time frame (e.g., a 4 hour period on the second Monday in February). All acceptances and rejections of offers occurred during this period. Keilin (2000) described the system as “problematic, subject to bottlenecks and gridlock, encouraging the violation of guidelines, and resulting in less-than-desirable outcomes for participants.”

In 1998-1999, the Association of Psychology Postdoctoral and Internship Centers (APPIC) switched to a system where applicants and internship sites were matched by computer. A major debate in this decision was whether a centralized system could handle the presence of couples. In the old decentralized scheme it was challenging for couples to coordinate their internship choices. Keilin (1998) reports that one partner could be put in the position of having to make an immediate decision about an offer without knowing the status of the other partner. Following the reforms of the National Resident Matching Program, a new scheme which allowed couples to jointly express their preferences was adopted.

With the permission of the APPIC, the company which runs the matching process, National

Matching Services, provided us with an anonymized dataset of the stated rank order lists of single doctors, couples and hospitals and hospital capacities for the first nine years of the centralized system. Because of privacy concerns, these data do not include any demographic information on applicants, and includes only limited identifying information on programs.

## 2.2 Stylized facts

This section identifies some stylized facts from the market for clinical psychologists. Other matching markets with couples appear to share many of features with this market, as we highlight below.

The data are the stated preferences of market participants, so their interpretation may require some caution. There are at least two parts to the process by which market participants form their preferences: (1) they determine which partners may be attractive and participate in interviews, and (2) after interviewing, they determine their rank ordering over partners. The model in this paper and the data do not allow us to say much about the first stage of the application process. In determining where to interview, applicants likely factor in the costs of traveling to interviews, the program's reputation and a host of other factors. Programs, among other things, consider the applicant's recommendation letters and suitability for their program in deciding whom to interview. Once market participants have learned about each other, they must come up with their rank ordering. For the empirical analysis in this section, we abstract away from the initial phase of mutual decisions of whom to interview, and our interpretation is that the data reflect the preferences that result *after* interviews.

Even with this interpretation, the reported post-interview preferences may be manipulations of the true post-interview preferences of market participants since truth-telling is not a dominant strategy for all market participants. However, there are at least two reasons why treating submitted preferences as true preferences may not be an unrealistic approximation. First, as noted in Section 6, the organizers of the APPIC match emphasize repeatedly that market participants should declare their preferences truthfully. Second, as we will see, Theorem 2 in Section 6 demonstrates that truth-telling is an approximate equilibrium in large markets.

Table 1 presents some summary statistics on the market. On average, per year, there are 3,010 single applicants and 19 pairs of applicants who participate as couples. In early years, there were just under 3,000 applicants, but the number of applicants has increased slightly in the most recent years. The number of applicants who participate as couples has remained relatively small, varying between 28 and 44, which is about 1% of all applicants.<sup>5</sup> In the National Resident Matching Program, from 1992-2009, there were on average 4.4% of applicants participating as couples, with slightly more

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<sup>5</sup>As the example in the next section shows, even one couple in the market may lead to non-existence of a stable matching.

couples participating in the most recent years (NRMP, 2009).

**Fact 1:** *Applicants who participate as couples constitute a small fraction of all participating applicants.*

Panel A of Table 1 shows the length of the rank order lists for applicants and programs. On average across years, single applicants rank between 7 and 8 programs. Since there are 1,094 programs on average, this means that the typical applicants ranks less than 1% of all possible programs. Even at the extreme, a single applicant ranks at most 6.7% of all possible programs. In the NRMP, the length of the applicant preference list is about 7-9 programs, which would be roughly 0.3% of all possible programs.<sup>6</sup> This may not be surprising because an applicant typically ranks a program only after she interviews at the program.

**Fact 2:** *The length of single applicants' rank order lists is small relative to the number of possible programs.*

Each entry in the rank order list of a couple is a pair of programs (or being unmatched). The typical rank order list of couples averages 81 program pairs. Since the rank order list of a couple has entries for both members, there are many duplicate programs, so we also report the number of distinct programs ranked by a couple. On average, there are 10 distinct programs listed by each couple member, which is larger than the average number of distinct programs listed by a single applicant. The maximum number of distinct programs ranked by a couple member is 1.9% of all programs.

**Fact 3:** *Applicants who participate as couples rank more programs than single applicants. However, the number of distinct programs ranked by a couple member is small relative to the number of possible programs.*

The next issue we examine is the distribution of applicant preferences. In Figure 1, we explore the popularity of programs in our data. For each program, we compute the total number of students who rank that program as their top choice. We order programs by this number, with the program with the highest number of top choices on the left and programs that no one ranks as their top choice on the right. Figure 1 shows the distribution of popularity for 2003. In this year, the most popular program was ranked as the top choice by 19 applicants, and there are 189 programs that are not ranked as a top choice by any applicant. The other years of our dataset display a similar pattern. Averaged across all years, the most popular program is ranked as a top choice by 24 applicants, and about 208 programs are not ranked as a top choice by anyone. The fraction of applicants ranking

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<sup>6</sup>This information is not available separately for single applicants and those who participate as couples in the NRMP.

the most popular program as their first choice is only 0.8%. (Recall that these are preferences stated after interviews have been conducted, so it does not preclude the possibility that there are popular programs that receive many applications but only interview a small subset of applicants.)

**Fact 4:** *The most popular programs are ranked as a top choice by a small number of applicants.*

The only identifying information we have on programs are geographic regions where they are located. The eleven geographic regions in our dataset are ten regions in the US, each of which corresponds to the first digit of the zip code of the program's location, and one region for all of Canada. Figure A1 illustrates these regions and shows the number of programs in each region. The Figure shows that programs are concentrated on the West Coast and in the Northeast. In Table 1, we report the number of distinct regions ranked by applicants. Half of single applicants rank at most two regions. Couples, on the other hand, tend to rank slightly more regions.

For a given couple rank order list, we also compute the fraction of entries on their submitted list that have both jobs in the same region. On average, 73% of a couple's rank order list is for programs in the same region.

**Fact 5:** *A pair of hospital programs ranked by doctors who participate as a couple tend to be in the same region.*

In the psychology market, there are about 1,100 residency programs. The average capacity is about 2.5 seats, and more than three quarters of programs have three or fewer spots. The total capacity of residency programs is smaller than the total number of applicants who participate, which implies that each year there will be unmatched applicants. This is also true in the NRMP where the number of positions per applicant ranges from 0.75-0.90 over the period 1995-2009 (NRMP, 2009).

Even though there are more doctors than programs, in the APPIC match, there are a sizable number of programs which are unfilled at the end of the regular match. According to the APPIC's statistics, during 1999-2007, on average 17% of programs had unfilled positions, and these were filled in a decentralized email aftermarket called the "Clearinghouse." In the NRMP, a similar number of programs had unfilled seats. In 2009, for instance, 12% of programs had unfilled positions, and these are filled in a an aftermarket managed by the NRMP called the "Scramble."

**Fact 6:** *Even though there are more applicants than positions, many programs still have unfilled positions at the end of the centralized match.*

### 2.3 Stable Matchings in the Market for Clinical Psychologists

We next investigate whether a stable matching exists in the psychologist market.<sup>7</sup> Roughly speaking, this is a matching such that there is no pair of hospital and applicant who prefer each other to the prescribed matching.<sup>8</sup> We use a variant of the procedure by Roth and Peranson (1999) to compute a stable matching.<sup>9</sup> For each of the 9 years of data, a stable matching exists in the market with couples. Table 2 shows that, averaged across years, more than a third of single applicants obtain their top choice averaged across each year in this stable matching, and 18% of couples obtain their top choice.<sup>10</sup> The number of unassigned couples is small (only about 5% of couples) while almost 20% of single applicants are unassigned.

**Fact 7:** *A stable matching exists in all nine years in the market for clinical psychologists.*

We also compare the assignment of single applicants at the stable matching we find in a market with couples to their assignment in the applicant-optimal stable matching in a market without couples in Table 2. While adding couples to the market could in principle affect the assignment received by many single applicants, in practice it has little effect. This can be seen by comparing the overall distribution of choice received for single applicant in a stable matching in markets without couples and with couples. Moreover, Table 3 reports the exact number of single applicants who receive a less preferred assignment in the market with couples. On average, there are 19 couples or 38 applicants who participate as couples in the market and because of their presence, only 63 single applicants obtain a lower choice. This corresponds to about 3 single applicants obtaining a different assignment per couple.

**Fact 8:** *Across stable matchings, most single applicants obtain the same position in the market without*

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<sup>7</sup>The model we analyze in this paper allows hospitals to have preferences over sets of doctors provided that the preferences are responsive. Our data on program rank order lists consist only of preferences over individual doctors. We do not know, for instance, whether a program prefers their first and fourth ranked applicants over their second and third ranked applicants. To compute a stable matching in the market for clinical psychologists, it is necessary to specify how comparisons between individual doctors relate to comparisons between sets of doctors. For the empirical computation, when comparing sets of doctors  $D_1$  and  $D_2$ , we assume that  $D_1$  is more preferred to  $D_2$  if the highest individually ranked doctor in  $D_1$  who is not in  $D_2$  is preferred to the highest individual ranked doctor in  $D_2$  who is not in  $D_1$ . This would imply that the first and fourth ranked applicant are preferred over the second and third ranked applicant.

<sup>8</sup>Section 3 provides a precise definition of our stability concept.

<sup>9</sup>Our variation has a different sequencing of applications from single applicants and couples than that described in Roth and Peranson (1999). That paper gives some evidence that these sequencing decisions have little impact on the success of the procedure.

<sup>10</sup>We focus on a particular stable matching in the market with couples, since we are unable to compute the entire set of stable matchings. There may be a reason to suspect that this set is small. In Table A.1 in the appendix, we compute stable matchings in the market without couples and find that very few applicants and programs have different assignments across the applicant-optimal and program-optimal stable matchings.

couples as in the market with couples.

In summary, this section describes some stylized facts that influence our choice of modeling assumptions. For the psychologist market, we are able to find a stable matching for each year for which we have data, which motivates our search for an existence result.

### 3 A Simple Theory of Matching with Couples

#### 3.1 Model

A matching market consists of hospitals, doctors, and their preferences. Let  $H$  be the set of hospitals plus an outside option  $\emptyset$  for doctors.  $S$  is the set of single doctors and  $C$  is the set of couples of doctors. Each couple is denoted by  $c = (f, m)$ , where  $f$  and  $m$  denote the first and second members of couple  $c$  respectively. When we need to refer to the members of a specific couple  $c$ , we sometimes write  $(f_c, m_c)$ . Let  $F = \{f | (f, m) \in C \text{ for some } m\}$  and  $M = \{m | (f, m) \in C \text{ for some } f\}$  be the sets of first and second members that form couples. Let  $D = S \cup F \cup M$  be the set of doctors.

Each single doctor  $s \in S$  has a preference relation  $R_s$  over  $H$ . We assume that preferences are strict: if  $hR_s h'$  and  $h'R_s h$ , then  $h = h'$ . We write  $hP_s h'$  if  $hR_s h'$  and  $h \neq h'$ . If  $hP_s \emptyset$ , we say that hospital  $h$  is **acceptable** to single doctor  $s$ .

Each couple  $c \in C$  has a preference relation  $R_c$  over  $H \times H$ , pairs of hospitals (and being unmatched). We assume that preferences of couples are strict with  $P_c$  denoting the asymmetric part of  $R_c$ . If  $(h, h')P_c(\emptyset, \emptyset)$ , then we say that pair  $(h, h')$  is acceptable to couple  $c$ . We say that hospital  $h$  is **listed by**  $R_c$  if there exists  $h' \in H$  such that either  $(h, h')P_c(\emptyset, \emptyset)$  or  $(h', h)P_c(\emptyset, \emptyset)$ .

Each hospital  $h \in H \setminus \{\emptyset\}$  has a preference relation over  $2^D$ , all possible subsets of doctors. We assume preferences of hospitals are strict. Let  $h \in H \setminus \{\emptyset\}$  and  $\kappa_h$  be a positive integer. We say that preference relation  $\succeq_h$  is **responsive with capacity**  $\kappa_h$  if it ranks a doctor independently of her colleagues and disprefers any set of doctors exceeding capacity  $\kappa_h$  to being unmatched (see Appendix A.1 for a formal definition). We follow much of the matching literature and assume that hospital preferences are responsive throughout the paper. Let  $R_h$  be the corresponding **preference list of hospital**  $h$ , which is the preference relation over individual doctors and  $\emptyset$ . We write  $dP_h d'$  if  $dR_h d'$  and  $d \neq d'$ . We say that doctor  $d$  is acceptable to hospital  $h$  if  $dP_h \emptyset$ . We write  $\succeq_H = (\succeq_h)_{h \in H}$ . We refer to a matching market  $\Gamma$  as a tuple  $(H, S, C, (\succeq_h)_{h \in H}, (R_i)_{i \in S \cup C})$ .

We proceed to define our stability concept in markets with couples. The descriptions are necessarily somewhat more involved than those in the existing literature because we allow for capacity of hospitals larger than one (we will elaborate on the issue in Section 3.1.1). First, it is convenient to introduce

the concept of hospital choices over permissible sets of doctors. For any set of doctors and couples  $D' \subseteq D \cup C$ , define

$$\begin{aligned} \mathcal{A}(D') = \{D'' \subseteq D \mid & \forall s \in S, \text{ if } s \in D'' \text{ then } s \in D', \\ & \forall c \in C, \text{ if } \{f_c, m_c\} \subseteq D'', \text{ then } (f_c, m_c) \in D', \\ & \text{if } f_c \in D'' \text{ and } m_c \notin D'', \text{ then } f_c \in D', \\ & \text{if } f_c \notin D'' \text{ and } m_c \in D'', \text{ then } m_c \in D'\}. \end{aligned}$$

In words,  $\mathcal{A}(D')$  is the collection of sets of doctors available for a hospital to employ when doctors (or couples of doctors)  $D'$  are applying to it. Underlying this definition is the distinction between applications by individual couple members and those by couples as a whole. For example, if  $(f, m) \in D' \cap C$  but  $f, m \notin D'$ , then the couple is happy to be matched to the hospital if and only if both members are employed together, while if  $(f, m) \notin D'$  but  $\{f, m\} \subseteq D'$ , then the couple is happy to have one member matched to the hospital but not together.

For any set  $D' \subseteq D \cup C$ , define the **choice** of hospital  $h$  given  $D'$ ,  $Ch_h(D')$ , to be the set such that

- $Ch_h(D') \in \mathcal{A}(D')$ ,
- $Ch_h(D') \succeq_h D''$  for all  $D'' \in \mathcal{A}(D')$ .

The choice  $Ch_h(D')$  is the most preferred subset of doctors among those in  $D'$  such that each couple is either chosen or not chosen together if they apply as a couple.<sup>11</sup>

A **matching** specifies which doctors are matched to which hospitals (if any). Formally, a matching  $\mu$  is a function defined on the set  $H \cup S \cup C$ , such that  $\mu(h) \subseteq D$  for every hospital  $h$ ,  $\mu(s) \in H$  for every single doctor  $s$ , and  $\mu(c) \in H \times H$  for every couple  $c$  where

- $\mu(s) = h$  if and only if  $s \in \mu(h)$  and
- $\mu(c) = (h, h')$  if and only if  $f_c \in \mu(h)$  and  $m_c \in \mu(h')$ .

When there are only single doctors in  $D'$ , the set  $\mathcal{A}(D')$  is simply the set of subsets of  $D'$ . Hence the choice  $Ch_h(D')$  is the subset of  $D'$  that is the most preferred by  $h$ . This is the standard definition of  $Ch_h(\cdot)$  in markets without couples (see Roth and Sotomayor (1990) for example), and hence the current definition is a generalization of the concept to markets with couples.

<sup>11</sup>We denote a singleton set  $\{x\}$  simply by  $x$  whenever there is no confusion.

A matching is **individually rational** if no player can be made better off by unilaterally rejecting some of the existing partners (see Appendix A.1 for a formal definition). We define different cases of a block as follows:

- (1) A pair of a single doctor  $s$  and a hospital  $h$  is a **block** of  $\mu$  if  $hP_s\mu(s)$  and  $s \in Ch_h(\mu(h) \cup s)$ .
- (2) (a) A coalition  $(c, h, h') \in C \times H \times H$  of a couple and two hospitals, where  $h \neq h'$ , is a **block** of  $\mu$  if
  - $(h, h')P_c\mu(c)$ ,
  - $f_c \in Ch_h(\mu(h) \cup f_c)$ , and
  - $m_c \in Ch_{h'}(\mu(h') \cup m_c)$ .
- (b) A pair  $(c, h) \in C \times H$  of a couple and a hospital is a **block** of  $\mu$  if
  - $(h, h)P_c\mu(c)$  and
  - $\{f_c, m_c\} \subseteq Ch_h(\mu(h) \cup c)$ .

A matching  $\mu$  is **stable** if it is individually rational and there is no block of  $\mu$ .

### 3.1.1 Discussion of the solution concepts

Models of matching with couples where hospitals have multiple positions are a particular form of many-to-many matching because each couple may seek two positions.<sup>12</sup> Various definitions of stability have been proposed for many-to-many matching, which differ based on the assumptions on what blocking coalitions are allowed (Sotomayor, 1999, 2004; Konishi and Ünver, 2006a; Echenique and Oviedo, 2006). Consequently, there are multiple possible stability concepts in matching with couples. The present definition of stability allows us to stay as close to the most commonly used pairwise stability as possible, by assuming away deviations involving large groups. Ruling out large coalitions appears to be reasonable because identifying and organizing large groups of agents may be difficult.

It is nevertheless important to understand whether our analysis is sensitive to a particular definition of stability. To address this issue, in Appendix A.2 we present an alternative definition of stability that allows for larger coalitions to block a matching. We show that all the results of this paper hold under that definition as well.

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<sup>12</sup>More precisely, Hatfield and Kojima (2008, 2009) point out that the model is subsumed by a many-to-many generalization of the matching model with contracts as analyzed by Hatfield and Milgrom (2005).

Most studies in matching with couples have focused on the case in which every hospital has capacity one.<sup>13</sup> Following the standard definition of stability in such models (see Klaus and Klijn (2005) for instance), we say that a matching  $\mu$  is **unit-capacity stable** if

- (1)  $\mu$  is individually rational,
- (2) there exists no single doctor-hospital pair  $s, h$  such that  $hP_s\mu(s)$  and  $sP_h\mu(h)$ , and
- (3) there exists no coalition by a couple  $c = (f, m) \in C$  and hospitals (or being unmatched)  $h, h' \in H$  with  $h \neq h'$  such that  $(h, h')P_c\mu(c)$ ,  $fR_h\mu(h)$  and  $mR_{h'}\mu(h')$ .<sup>14</sup>

Our concept of stability is equivalent to the unit-capacity stability as defined above if every hospital has responsive preferences with capacity one. To see this, first observe that condition (3) of unit-capacity stability is equivalent to the nonexistence of a block as defined in condition (2a) of our stability concept. Moreover, condition (2b) of our stability concept is irrelevant when each hospital has capacity one because a hospital with capacity one never prefers to match with two members of a couple. Finally, the remaining conditions for unit-capacity stability have direct counterparts in our definition of stability. Thus the stability concept employed in this paper is a generalization of the standard concept to the case where hospitals have multiple positions.

Also note that our stability concept is equivalent to the standard definition of (pairwise) stability when there exist no couples. More specifically, condition (2) of our stability concept is irrelevant if couples are not present, and condition (1) is equivalent to the nonexistence of a blocking pair which, together with individual rationality, defines stability in markets without couples.

### 3.2 The Problem with Couples

We illustrate how the existence of couples poses problems in the theory of two sided matching. To understand the role of couples, however, it is useful to start by considering a matching without couples. In that context, the (doctor-proposing) **deferred acceptance algorithm** defined below always produces a stable matching (Gale and Shapley, 1962).

**Algorithm 1.** DOCTOR-PROPOSING DEFERRED ACCEPTANCE ALGORITHM

Input: a matching market  $(H, S, (\succeq_h)_{h \in H}, (R_s)_{s \in S})$  without couples.

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<sup>13</sup>Some papers consider multiple positions of hospitals but treat a hospital with capacity larger than one as multiple hospitals with capacity one each. This approach is customary and innocuous when there exists no couple because most stability concepts are known to coincide in that setting. However the approach has a consequence if couples exist since it leads to a particular stability concept. A different modeling approach is pursued by McDermid and Manlove (2009).

<sup>14</sup>We adopt the notational convention that  $dR_{\emptyset}d'$  for any  $d, d' \in D \cup \emptyset$ .

- Step 1: Each single doctor applies to her first choice hospital. Each hospital rejects its least-preferred doctor in excess of its capacity and all unacceptable doctors among those who applied to it, keeping the rest of the doctors temporarily (so doctors not rejected at this step may be rejected in later steps).

In general,

- Step  $t$ : Each doctor who was rejected in Step  $(t-1)$  applies to her next highest choice (if any). Each hospital considers these doctors *and* doctors who are temporarily held from the previous step together, and rejects the least-preferred doctors in excess of its capacity and all unacceptable doctors, keeping the rest of the doctors temporarily (so doctors not rejected at this step may be rejected in later steps).

The algorithm terminates at a step where no doctor is rejected. The algorithm always terminates in a finite number of steps. At that point, all tentative matchings become final. Gale and Shapley (1962) show that for any given market without couples, the matching produced by the deferred acceptance algorithm is stable. Furthermore, they show that it is the doctor-optimal stable matching, the stable matching that is weakly preferred to any other stable matching by all doctors.

By contrast, stable matchings do not necessarily exist when there exists a couple in the market (shown by Roth (1984) and an unpublished work by Sotomayor). This fact is illustrated in the following example, based on Klaus and Klijn (2005).

**Example 1.** Let there be a single doctor  $s$  and a couple  $c = (f, m)$  as well as two hospitals  $h_1$  and  $h_2$ , each with capacity one. Suppose the acceptable matches for each agent, in order of preference, are given by:

$$\begin{array}{ll} R_c : (h_1, h_2) & R_s : h_1, h_2 \\ \succeq_{h_1} : f, s & \succeq_{h_2} : s, m. \end{array}$$

We illustrate that there is no stable matching in this market, by considering each possible matching.

- (1) Suppose  $\mu(c) = (h_1, h_2)$ . Then single doctor  $s$  is unmatched. Thus single doctor  $s$  and hospital  $h_2$  block  $\mu$  because  $s$  prefers  $h_2$  to her match  $\mu(s) = \emptyset$  and  $h_2$  prefers  $s$  to its match  $\mu(h_2) = m$ .
- (2) Suppose  $\mu(c) = (\emptyset, \emptyset)$ .

- (a) If  $\mu(s) = h_1$ , then  $(c, h_1, h_2)$  blocks  $\mu$  since couple  $c$  prefers  $(h_1, h_2)$  to their match  $\mu(c) = (\emptyset, \emptyset)$ , hospital  $h_1$  prefers  $f$  to its match  $\mu(h_1) = s$  and hospital  $h_2$  prefers  $m$  to its match  $\mu(h_2) = \emptyset$ .
- (b) If  $\mu(s) = h_2$  or  $\mu(s) = \emptyset$ , then  $(s, h_1)$  blocks  $\mu$  since single doctor  $s$  prefers his first choice hospital  $h_1$  to both hospital  $h_2$  and  $\emptyset$  while  $h_1$  prefers  $s$  to its match  $\mu(h_1) = \emptyset$ .

Klaus and Klijn (2005) identify a sufficient condition to guarantee the existence of a stable matching called weak responsiveness. A couple's preferences are said to be responsive if an improvement in one couple member's assignment is an improvement for the couple. Preferences are said to be weakly responsive if the requirement applies to all acceptable positions.<sup>15</sup> The preferences of couples in Example 1 do not satisfy this condition. If, for instance, the couple's preferences are  $(h_1, h_2), (h_1, \emptyset), (\emptyset, h_2), (\emptyset, \emptyset)$ , in order of preference, then it satisfies responsiveness and a stable matching exists. Klaus and Klijn (2005) write that "responsiveness essentially excludes complementarities in couples' preferences." They showed that:

- (1) if the preferences of every couple are weakly responsive, then there exists a stable matching.
- (2) if there is at least one couple whose preferences violate weak responsiveness while satisfying a condition called "restricted strict unemployment aversion," then there exists a preference profile of other agents such that preferences of all other couples are weakly responsive but there exists no stable matching.

Their second result says that the class of weakly responsive preferences is the "maximal domain" of preferences. That is, it is the weakest possible condition that can be imposed on individual couples' preferences that guarantees the existence of stable matchings.<sup>16</sup>

There seem to be many situations in which couple preferences violate weak responsiveness. One reason may be geographic, as stated as Fact 5 in Section 2.2: both programs ranked as a pair by a couple tend to be in the same geographic region. For example, the first choice of a couple of medical residents may be two residency programs in Boston and the second may be two programs in Los Angeles, while one member working in Boston and the other working in Los Angeles could be unacceptable because these two cities are too far away from each other. The coordinator of the Association of Psychology Postdoctoral and Internship Centers (APPIC) matching program writes in Keilin (1998) that "most couples want to coordinate their internship placements, particularly with

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<sup>15</sup>See Klaus et al. (2009) for formal definition.

<sup>16</sup>Hatfield and Kominers (2009) show that the substitutes condition is a maximal domain in the absence of restricted strict unemployment aversion.

regard to geographic location.” This suggests that violation of weak responsiveness due to geographic preferences is one of the representative features of couple preferences.

To further study this question empirically, we analyze the data on the stated preferences of couples from the APPIC.<sup>17</sup> During years for which we have data (1999–2007), preferences of only one couple out of 167 satisfy weak responsiveness. Thus the data suggest, in light of the results of Klaus and Klijn (2005), that it is virtually impossible to guarantee the existence of a stable matching in such markets with couples based on a domain restriction of preferences.

However, the fact that the preferences of the overwhelming majority (166 out of 167) of couples violate weak responsiveness does not mean that a stable matching does not exist in the psychologist market. Stable matchings have been found in many labor markets despite the presence of couples, and as we described in Section 2.3, we find a stable match for each of the nine years of the psychology market for which we have data. This motivates our desire to understand what market features enable the existence of stable matchings most of the time, when the known sufficient conditions on couples’ preferences do not guarantee existence.

### 3.3 Sequential Couples Algorithm

The original deferred acceptance algorithm does not incorporate applications by couples. We consider an extension of the algorithm, which we call the **sequential couples algorithm**. While we defer a formal definition to Appendix A.3 for expositional simplicity, we offer an informal description as follows.

- (1) run a deferred acceptance algorithm for a sub-market composed of all hospitals and single doctors, but without couples,
- (2) one by one, place couples by allowing each couple to apply to pairs of hospitals in order of their preferences (possibly displacing some doctors from their tentative matches), and
- (3) one by one, place singles who were displaced by couples by allowing each of them to apply to a hospital in order of her preferences.

We say that the sequential couples algorithm **succeeds** if there is no instance in the algorithm in which an application is made to a hospital where an application has previously been made by a member (or both members) of a couple except for the couple who is currently applying. Otherwise, we declare a failure and terminate the algorithm.

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<sup>17</sup>Since truth-telling is not necessarily a dominant strategy for couples, the use of stated preferences is potentially problematic. However, Theorem 2 in this paper provides formal defense for this assumption by demonstrating that truth-telling is an approximate equilibrium in large markets.

In practice, a matching clearinghouse would be unlikely to declare failure when the sequential couples algorithm fails, and instead consider some procedure to try to assign the remaining couples. Following the sequential couples algorithm with such a procedure is the main idea behind the Roth-Peranson algorithm (Roth and Peranson, 1999), which is the basis for the mechanism used in the NRMP, APPIC, and other labor markets. The Roth-Peranson algorithm first runs a sequential couples algorithm. If the latter algorithm succeeds, then the Roth-Peranson algorithm produces the matching reached by the sequential couples algorithm. If the sequential couples algorithm fails, then the Roth-Peranson algorithm proceeds to a second stage, in which doctors who are potentially members of blocking pairs can apply again. In each of the subsequent steps the algorithm allows agents to form blocking pairs, eliminating instances of instability one by one, in a manner similar to Roth and Vate (1990). Since a stable matching does not necessarily exist in markets with couples, the Roth-Peranson algorithm could cycle without terminating. The algorithm forces termination in such a case without finding a stable matching. The complete description of the algorithm, specifically how the algorithm terminates when the algorithm does not find a stable matching, is not publicly available, but a more detailed description than the one provided here is offered by Roth and Peranson (1999).

Despite the similarity, the sequential couples algorithm and the Roth-Peranson algorithm are different in two aspects. First, the Roth-Peranson algorithm proceeds with the algorithm trying to find a stable matching even in cases when a failure is declared in the sequential couples algorithm. Second, in the Roth-Peranson algorithm, when a couple is added to the market with single doctors, any single doctor who is displaced by the couple is placed before another couple is added. By contrast, the sequential couples algorithm holds any displaced single doctor without letting her apply, until it processes applications by all couples.<sup>18</sup>

The reason we focus on this simplified procedure is that the success of the sequential couples algorithm turns out to play an important role as the next proposition relates it to the existence of a stable matching (the proof is in Appendix A.3).

**Lemma 1.** *If the sequential couples algorithm succeeds, then the resulting matching is stable.*

To illustrate the main idea of Lemma 1, we consider how the sequential couples algorithm proceeds for the market in Example 1. In Step 1 of the algorithm, we run the doctor-proposing deferred acceptance algorithm in the sub-market without couples. Single doctor  $s$  proposes to hospital  $h_1$  and is assigned there. Then in Step 2, we let couple  $c$  apply to their top choice  $(h_1, h_2)$ . Couple member  $f$  is preferred to  $s$  by  $h_1$  and couple member  $m$  is preferred to a vacant position by  $h_2$ . Thus  $f$  and  $m$  are tentatively assigned to  $h_1$  and  $h_2$  respectively while  $s$  is rejected. Then in Step 3, we let  $s$  apply to

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<sup>18</sup>As we will point out subsequently, our result also holds if we follow the sequencing of the Roth-Peranson algorithm. We chose the current definition of the sequential couples algorithm for expositional simplicity.

her next highest choice. In this case, she applies to hospital  $h_2$ , where a couple member  $m$  has applied and been assigned before. At this point we terminate the algorithm and declare that it has failed.

To see why declaring a failure of the sequential couples algorithm is useful, suppose that we hypothetically continue the algorithm by allowing  $h_2$  reject  $m$  as  $h_2$  prefers  $s$  to  $m$ . Then the couple prefers being unassigned rather than having only  $f$  be matched to  $h_1$ , so doctor  $f$  would like to withdraw his assignment from hospital  $h_1$ . Suppose we terminate the algorithm at this point once  $f$  becomes unmatched. Then the resulting matching assigns no doctor to  $h_1$  and  $s$  to  $h_2$ . This matching is unstable because doctor  $s$  can block with hospital  $h_1$ . On the other hand, if we continue the algorithm further by allowing  $s$  to match with  $h_1$ , then the resulting matching is identical to the one obtained at the end of Step 1 of the sequential couples algorithm. This suggests that reasonable algorithms would cycle without terminating in this market.

The idea of declaring failure of the sequential couples algorithm is to avoid a situation like the above example, and turns out to be a useful criterion for judging whether the algorithm produces a stable matching. Of course, the algorithm sometimes fails even if there exists a stable matching, so the success of the algorithm is only a sufficient condition for the existence of a stable matching. What is remarkable is that looking at this particular sufficient condition turns out to be enough for establishing that a stable matching exists with a high probability in the environment we study in this paper. Moreover, there is a sense in which it is *necessary* to use an algorithm that finds a stable matching only in some instances, rather than one that always finds a stable matching whenever it exists. Ronn (1990) shows that the problem of determining whether a market with couples has a stable matching or not is computationally hard (NP-complete). The result suggests that it may be inevitable to employ an approach that does not always find a stable matching like our sequential couples algorithm.

Example 1 illustrates that the sequential couples algorithm does not necessarily succeed, and suggests that any market of finite size would allow such a failure. We instead consider a large market environment with a random component in the preferences of the market participants. Our contribution is to demonstrate that, with high probability, the sequential couples algorithm succeeds, and hence a stable matching exists in this environment.

## 4 Large Markets

### 4.1 Random Markets

We have seen that a stable matching does not necessarily exist in a finite matching market with couples. To investigate how often a stable matching exists in large market, we introduce the following

random environment. A **random market** is a tuple  $\tilde{\Gamma} = (H, S, C, \succeq_H, k, \mathcal{P}, \mathcal{Q}, \rho)$ , where  $k$  is a positive integer,  $\mathcal{P} = (p_h)_{h \in H}$  and  $\mathcal{Q} = (q_h)_{h \in H}$  are probability distributions on  $H$ , and  $\rho$  is a function which maps two preferences over  $H$  to a preference list for couples (explained below). Each random market induces a market by randomly generating preferences of doctors as follows:

**Preferences for Single Doctors:** For each single doctor  $s \in S$ ,

- Step 1: Select a hospital independently from distribution  $\mathcal{P}$ . List this hospital as the top ranked hospital of single doctor  $s$ .

In general,

- Step  $t \leq k$ : Select a hospital independently from distribution  $\mathcal{P}$  until a hospital is drawn that has not been previously drawn in steps 1 through  $t - 1$ . List this hospital as the  $t$ th most preferred hospital of single doctor  $s$ .

Single doctor  $s$  finds these  $k$  hospitals acceptable, and all other hospitals unacceptable. For example, if  $\mathcal{P}$  is the uniform distribution on  $H$ , then the preference list is drawn from the uniform distribution over the set of all preference lists of length  $k$ .

**Preferences for Doctors who are Couples:** Couples' preferences are formed by drawing preferences,  $R_f$  and  $R_m$ , for each doctor in the couple  $c = (f, m)$ .  $R_f$  is constructed from the same process used to generate preferences for a single doctor, except that the hospitals are drawn from distribution  $\mathcal{Q}$  instead of  $\mathcal{P}$ . Likewise,  $R_m$  is generated using  $\mathcal{Q}$ .

To construct the preference list for the couple  $c = (f, m)$ , define  $\rho(R_f, R_m)$  to be a preference of the couple with the following restriction: if  $(h_1, h_2)$  is acceptable according to  $\rho(R_f, R_m)$ , then  $h_1 R_f \emptyset$  and  $h_2 R_m \emptyset$ . This is the only restriction we place on  $\rho$ .

**Preferences for Hospitals:** Each hospital  $h$  has a responsive preference relation defined over sets of doctors  $\succeq_h$  such that all doctors are acceptable. The preference list-capacity pair consistent with  $\succeq_h$  is denoted by  $(R_h, \kappa_h)$ .

## Discussion of modeling choices

We are specializing the structure of the model in several important ways. One important modeling choice is that doctor preferences are drawn independently from one another, and the way in which each doctor's preference list is drawn also follows a particular procedure. While this framework excludes

some cases, it has been used in several papers on matching such as Immorlica and Mahdian (2005), Kojima and Pathak (2009), and Manea (2009). One cannot dispense with these restrictions completely, as some of our results fail when these assumptions are violated. For example, Section 6 establishes an approximate incentive compatibility of a class of mechanisms in large matching markets, but the result fails under preference distributions violating our assumption.<sup>19</sup>

We allow  $\mathcal{Q}$  for couples to be different from  $\mathcal{P}$  for single doctors, sacrificing simplicity. We chose to do so because couples in practice could have very different views on desirability of hospitals from those held by single doctors.

The function  $\rho$  is a mapping that outputs a preference relation for each couple  $(f, m)$  given the pair of preferences  $R_f$  and  $R_m$  over  $H$ . One could interpret  $\rho(R_f, R_m)$  as describing the outcome of household bargaining when preferences of the members are  $R_f$  and  $R_m$ , respectively. For example, the function  $\rho$  can represent a process in which any pair of hospitals that are too far away from each other is declared unacceptable, which seems to be consistent with the observed rank order lists of couples described earlier. We remain agnostic about  $\rho$  except that a hospital pair  $(h, h')$  is weakly acceptable for the couple under  $\rho(R_f, R_m)$  only when  $h$  and  $h'$  are listed under  $R_f$  and  $R_m$ , respectively. In other words, no hospital appears in the preference list of a couple unless it is considered by the relevant member of the couple. All our results are unchanged if we allow the function  $\rho$  to vary across different couples, but we model a common function  $\rho$  for all couples for expositional simplicity.

Some NRMP participants who participate as couples are advised to form preferences by first forming individual rank order lists after interviewing with programs. Then, these individual's lists serve as an input into the joint ranking of the couple. For instance, medical students who are couples at the University of Kansas Medical School are suggested to make a list of all possible program pair combinations from both individual rank order lists by computing the difference between the ranking number of the program on each individual's rank order list and trying to minimize this difference in their joint rank order list. This would be one example of a  $\rho$  function.<sup>20</sup>

The probabilistic structure we place on doctor preferences is unneeded for hospital preferences. Rather, hospital preferences can be arbitrary except for two important restrictions. First, hospital preferences are assumed to be responsive as in much of the literature on two-sided matching. The

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<sup>19</sup>There is an example in Immorlica and Mahdian (2005) where preference distributions violate our assumption and the result fails even without couples. Since the current model is a generalization of theirs, the counterexample applies. See also the discussion in Kojima and Pathak (2009).

<sup>20</sup>The details on this advice are available at <http://www.kumc.edu/som/medsos/cm.html>, accessed on March 20, 2010. The clearinghouse for new doctors in Scotland only allows couple members to submit individual rank order lists, in contrast to the model we analyze here. In that context, their mechanism combines these lists into a preference over pairs for the couple using their individual lists and a table of positions that are determined to be geographically compatible by the mechanism. See the discussion of the Scottish Foundation Allocation Scheme at <http://www.nes.scot.nhs.uk/sfas/About/default.asp>, accessed on March 29, 2010.

labor market clearinghouses which motivate our study impose this restriction by eliciting preferences over individual doctors. The second important assumption on hospital preferences is that hospitals find all doctors acceptable. We make this assumption so that there are enough hospitals that can actually hire doctors in large markets. At first glance, this assumption seems violated in the data from the market for clinical psychologists as no program submits a rank order list of all doctors. In our data, however, the vast majority of programs rank every doctor who has ranked them, which might suggest that most applicants would in fact be acceptable to a program had they interviewed there. The results follow, at additional notational complexity, in a model where many, but not all, hospitals find all doctors acceptable.

## 4.2 Regular Sequence of Random Markets

To analyze limit behavior of the matching market as the market becomes large, we consider a sequence of markets of different sizes. A **sequence of random markets** is denoted by  $(\tilde{\Gamma}^1, \tilde{\Gamma}^2, \dots)$ , where  $\tilde{\Gamma}^n = (H^n, S^n, C^n, \succeq_{H^n}, k^n, \mathcal{P}^n, \mathcal{Q}^n, \rho^n)$  is a random market in which  $|H^n| = n$  is the number of hospitals. Consider the following regularity conditions.

**Definition 1.** A sequence of random markets  $(\tilde{\Gamma}^1, \tilde{\Gamma}^2, \dots)$  is **regular** if there exist  $\lambda > 0$ ,  $a \in [0, \frac{1}{2})$ ,  $b > 0$ ,  $r \geq 1$ , and positive integers  $k$  and  $\bar{\kappa}$  such that for all  $n$ ,

- (1)  $k^n = k$ ,
- (2)  $|S^n| \leq \lambda n$ ,  $|C^n| \leq bn^a$ ,
- (3)  $\kappa_h \leq \bar{\kappa}$  for all hospitals  $h$  in  $H^n$ ,
- (4)  $\frac{p_h}{p_{h'}} \in [\frac{1}{r}, r]$  and  $\frac{q_h}{q_{h'}} \in [\frac{1}{r}, r]$  for all hospitals  $h, h'$  in  $H^n$ .

Condition (1) assumes that the length of doctors' preference lists does not grow when the number of market participants grow. This assumption is motivated by Facts 2 and 3 in Section 2.2 that the length of single doctors' preference lists is small relative to the number of hospitals. Condition (2) requires that the number of single doctors does not grow much faster than the number of hospitals. Moreover, couples do not grow at the same rate as the number of hospitals and instead grow at the slower rate of  $O(n^a)$  where  $a \in [0, \frac{1}{2})$ . This condition is motivated by Fact 1 that the number of couples is small compared with the number of hospitals or single doctors. Condition (3) requires that the capacity of each hospital is bounded. This condition is not needed for the existence result of a stable matching, and we use it only for our incentive result. Condition (4) requires that the popularity

of different hospitals (as measured by the probability of being listed by doctors as acceptable) does not vary too much, as suggested by Fact 4. Allowing lengths of preference lists to be different from doctor to doctor does not change any of our results, as long as there is an upper bound  $k$  of list lengths where  $k$  is a constant independent of  $n$ .<sup>21</sup>

This paper focuses on regular sequences of random markets and makes use of each condition in our arguments.

## 5 Existence of Stable Matchings

As seen in Example 1, a stable matching does not necessarily exist when some doctors are couples. However, there is a sense in which a stable matching is likely to exist if the market is large. This claim is formalized in the following result on asymptotic existence for a regular sequence of random markets (Definition 1).

**Theorem 1.** *Suppose that  $(\tilde{\Gamma}^1, \tilde{\Gamma}^2, \dots)$  is a regular sequence of random markets. Then the probability that there exists a stable matching in the market induced by  $\tilde{\Gamma}^n$  converges to one as the number of hospitals  $n$  approaches infinity.*

We defer the formal proof to Appendix A.3 and describe the argument here. Our proof involves analysis of the sequential couples algorithm in a regular sequence of random markets. By Lemma 1, we know that a stable matching exists whenever the algorithm succeeds. Our proof strategy is to show that the probability that the sequential couples algorithm succeeds converges to one as the market size approaches infinity.

Suppose that there are a large number of hospitals in the market. Given our assumptions on the distribution of couples' preferences, different couples are likely to prefer different pairs of hospitals. Hence, in Step 2 of the algorithm, members of two distinct couples are unlikely to apply to the same hospital. In such an instance, this step of the algorithm tentatively places couples without terminating. Given that, it suffices to show that the single doctors displaced in Steps 2 and 3 (if any) are likely to be placed without applying to a hospital where a couple has applied. To show this, first we demonstrate that if the market is large, then it is a high probability event that there are a large number of hospitals with vacant positions at the end of Step 2 (even though there could be more applicants than positions).<sup>22</sup> Then, any single doctor is much more likely to apply to a hospital

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<sup>21</sup>Fact 3 indicates that applicants who participate as couples tend to rank more programs than applicants who participate as singles, but both sets of applicants' rank order lists are small relative to the number of programs.

<sup>22</sup>Note that the feature that there are many hospitals with vacant positions is consistent with Fact 6 in Section 2.2, which states that there are many resident programs with vacant positions in practical matching markets.

with a vacant position than to one of the hospitals that has already received an application by a couple member. Since every doctor is acceptable to any hospital by assumption, a doctor is accepted whenever an application is made to a vacant position. With high probability the algorithm places all the single doctors in Step 3, resulting in a success. Together with Lemma 1, we conclude that if the market is large enough, then the probability that there exists no stable matching can be made arbitrarily small. This completes the argument.

As explained in Section 3.3, the sequential couples algorithm is similar to but slightly different from the Roth-Peranson algorithm in the order of which doctors apply to hospitals. However, it is clear from the proof that the argument can be modified for the Roth-Peranson algorithm. Therefore, we have the following result as a corollary.

**Corollary 1.** *Suppose that  $(\tilde{\Gamma}^1, \tilde{\Gamma}^2, \dots)$  is a regular sequence of random markets. Then the probability that the Roth-Peranson algorithm produces a stable matching in the market induced by  $\tilde{\Gamma}^n$  converges to one as the number of hospitals  $n$  approaches infinity.*

In Appendix A.3, we show that if the number of couples is bounded along the sequence (that is,  $a = 0$  in Definition 1), then the probability that there is no stable matching approaches zero at least with the rate of convergence  $O(1/n)$ .

## 6 Incentives

The last section establishes that the sequential couples algorithm finds a stable matching with a high probability in large markets with respect to reported preferences that follow certain distributional assumptions. In practice, however, preferences are private information of market participants, and the matching clearinghouse needs to elicit the information from them. Thus a natural question is whether there is a mechanism that induces participants to report true preferences and produces a stable matching with respect to the true preferences. This problem is addressed in this Section.

One motivation for studying this question comes from the market for clinical psychologists. The following advice is given to participants:<sup>23</sup>

IMPORTANT: There is only one correct “strategy” for developing your Rank Order List: simply list your sites based on your true preferences, without consideration for where you

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<sup>23</sup>“FAQ for Internship Applicants” in the APPIC website, <http://www.appic.org/match/5.2.1.2.6.html>, accessed on 11/11/2009.

believe you might be ranked by them. List the site that you want most as your #1 choice, followed by your next most-preferred site, and so on.

The previous paragraph is so important that we are going to repeat it: simply list your sites based on your true preferences.

Similar recommendations are made in other labor markets with couples. Below is the advice for participants offered by the National Resident Matching Program (NRMP).<sup>24</sup>

Programs should be ranked in sequence, according to the applicant's true preferences. . . . It is highly unlikely that either applicants or programs will be able to influence the outcome of the Match in their favor by submitting a list that differs from their true preferences.

In these quotes, market participants are advised to report their true preferences to the matching authority, even though no existing study analyzes formally when truth-telling is optimal in markets with couples.

Motivated by these observations, we study incentives for manipulation in a market with couples. At a first glance, a positive result seems elusive: there exists no mechanism that is stable and strategy-proof even without couples (Roth, 1982). We seek, therefore, an idea of approximate incentive compatibility in the context of large matching markets.

## 6.1 Mechanism

To consider the incentives for manipulation in a market with couples, we consider a mechanism which builds on the sequential couples algorithm. We denote this mechanism by  $\varphi$ . The reason we focus on mechanism  $\varphi$  is that we need to specify what matching is returned by the mechanism when the sequential couples algorithm fails. We defer a formal definition of the mechanism to the Appendix and present an informal description here.

- (1) run a deferred acceptance algorithm for a sub-market composed of all hospitals and single doctors, but without couples,
- (2) one by one, place couples by allowing each couple to apply to pairs of hospitals in order of their preferences (possibly displacing some doctors from their tentative matches), and
- (3) one by one, place singles who were displaced by couples by allowing each of them to apply to a hospital in order of her preferences.

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<sup>24</sup>“Rank Order Lists” in the NRMP website, [http://www.nrmp.org/fellow/rank\\_order.html](http://www.nrmp.org/fellow/rank_order.html), accessed on 11/11/2009.

This algorithm terminates in a finite number of steps. We define  $\varphi$  to be the mechanism that, given all reported preferences and capacities, produces the matching reached at the end of this algorithm. This mechanism is analogous to the sequential couples algorithm. In fact, the algorithm defining  $\varphi$  proceeds identically to the sequential couples algorithm as long as the latter succeeds. Unlike the sequential couples algorithm, however, in mechanism  $\varphi$  we do not declare failure when someone applies to a hospital to which a couple member has already applied. Instead, we allow the new applicant to be assigned to the hospital. If the new applicant applies to a hospital in which a couple member is already tentatively matched, then we allow the applicant to displace the couple member. If a couple member is displaced, then we assume that the other member of that couple withdraws application from the current match (if any), and the couple applies to their next preferred hospital pair. More specifically, the algorithm forces each (single or couple) doctor to apply from the top-ranked hospital (pair) and prevents her from applying again to the same hospital (pair). By construction this algorithm terminates in a finite number of steps, at which point the tentative matching becomes final. This mechanism does not necessarily produce a stable matching. However, since this algorithm coincides with the sequential couples algorithm whenever the latter succeeds, the proof of Theorem 1 implies that the probability that  $\varphi$  produces a stable matching converges to one as the market size becomes infinitely large.

## 6.2 Equilibrium

To study incentives of participants in mechanism  $\varphi$ , we consider a Bayesian game in which both hospitals and doctors report their preferences strategically to the matching authority. Kojima and Pathak (2009) study a similar model without couples, but the current analysis involves a number of additional considerations due to the existence of couples. For instance, if there are no couples, then reporting true preferences is a dominant strategy for every doctor in  $\varphi$ , but such a result is no longer true if there are couples.<sup>25</sup> As a result, we need to analyze strategic behavior by all market participants including doctors, rather than only hospitals as in Kojima and Pathak (2009).

A **matching game** is a Bayesian game specified by  $\hat{\Gamma} = (H, S, C, (\mathcal{U}_i, F_i)_{i \in H \cup S \cup C})$ . The set of players is  $H \cup S \cup C$ , the set of all hospitals and doctors (including both singles and couples). For each player  $i$ , the set  $\mathcal{U}_i$  represents the set of utility types for  $i$ , with each element  $u_i$  specifying a utility function for  $i$ .  $F_i$  is a probability distribution over  $\mathcal{U}_i$ . Types  $(u_i)_i$  are independently distributed across agents.

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<sup>25</sup>To see why truth-telling is a dominant strategy for each doctor if there are no couples, note that  $\varphi$  is equivalent to the doctor-proposing deferred acceptance algorithm if there are no couples. Truth-telling is a dominant strategy for every doctor under the doctor-proposing deferred acceptance algorithm (Dubins and Freedman, 1981; Roth, 1982), thus the assertion follows.

All the players move simultaneously. At strategy  $\sigma_i$ , a player  $i$  submits an ordinal preference relation  $\sigma_i(u_i)$  upon observing her own type  $u_i$ , but not the realized types of the other players. We assume that hospitals are allowed to submit only responsive preferences and single doctor and couples can submit any preference relation in Section 3.1. Once all players report their preferences, each player  $i$  receives a matching resulting from  $\varphi$  under the submitted preferences.<sup>26</sup>

Given  $\varepsilon \geq 0$ , a strategy profile  $\sigma^*$  is an  $\varepsilon$ -**Bayes Nash equilibrium** if there exists no  $i \in H \cup S \cup C$ ,  $u_i \in \mathcal{U}_i$  and strategy  $\sigma_i$  such that

$$E[u_i(\varphi_i(\sigma_i(u_i), \sigma_{-i}^*(u_{-i})))] > E[u_i(\varphi_i(\sigma^*(u)))] + \varepsilon,$$

where  $\varphi_i(\sigma_i(u_i), \sigma_{-i}^*(u_{-i}))$  and  $\varphi_i(\sigma^*(u))$  are the matchings for  $i$  when reported preference profiles are  $(\sigma_i(u_i), (\sigma_j^*(u_j))_{j \neq i})$ , and  $(\sigma_j^*(u_j))_{j \in H \cup S \cup C}$  respectively. That is, a strategy profile is an  $\varepsilon$ -Bayes Nash equilibrium if no player of any type can gain utility of more than  $\varepsilon$  by unilateral deviation. This concept is a generalization of the standard Bayes-Nash equilibrium and coincides with it if  $\varepsilon = 0$ . We say that a strategy profile  $\sigma$  is **truth-telling** if for every  $i$  and type  $u_i$ , strategy  $\sigma_i(u_i)$  is the ordinal preference represented by  $u_i$ . That is, a strategy profile is truth-telling if every player of any type reports their true ordinal preferences.

To analyze incentive compatibility in large markets, consider a sequence  $(\hat{\Gamma}^1, \hat{\Gamma}^2, \dots)$  of matching games, where  $\hat{\Gamma}^n = (H^n, S^n, C^n, (\mathcal{U}_i, F_i)_{i \in H^n \cup S^n \cup C^n})$  is the game in which there are  $n$  hospitals. Normalize utility functions such that utility of being unmatched is zero for every player. We consider the following definition of regularity for a sequence of matching games:

**Definition 2.** A sequence of matching games  $(\hat{\Gamma}^1, \hat{\Gamma}^2, \dots)$  is **regular** if there exist  $\lambda > 0$ ,  $a \in [0, \frac{1}{2})$ ,  $b > 0$ ,  $r \geq 1$ , positive integers  $k$  and  $\bar{\kappa}$ , and distributions  $\mathcal{P}^1, \mathcal{Q}^1, \mathcal{P}^2, \mathcal{Q}^2, \dots$  such that for all  $n$ ,

- (1) Distribution  $F_i$  for  $i \in S^n$  (respectively  $i \in C^n$ ) is such that the distribution on ordinal preferences  $R_i$  represented by the utility function  $u_i$  follows the processes described in Section 4.1 associated with  $\mathcal{P}^n$  (respectively  $\mathcal{Q}^n$  and  $\rho^n$ ) and  $k$ .
- (2)  $|S^n| \leq \lambda n$ ,  $|C^n| \leq bn^a$ ,
- (3) For each  $n$ ,  $h \in H^n$ , ordinal preferences of  $h$  represented by each type  $u_h$  satisfy assumptions in Section 4.1 representing a responsive preference with capacity at most  $\bar{\kappa}$ ,
- (4)  $\frac{p_h}{p_{h'}} \in [\frac{1}{r}, r]$  and  $\frac{q_h}{q_{h'}} \in [\frac{1}{r}, r]$  for all hospitals  $h, h'$  in  $H^n$ .

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<sup>26</sup>Thus if player  $i$ 's type is  $u_i$  and matching  $\mu$  results, then she receives utility  $u_i(\mu(i))$ .

$$(5) \sup_{n \in \mathbb{N}, D^n, h \in H^n, u_h \in \mathcal{U}_h} u_h(D'), \sup_{n \in \mathbb{N}, s \in S^n, h \in H^n, u_s \in \mathcal{U}_s} u_s(h), \sup_{n \in \mathbb{N}, c \in C^n, h, h' \in H^n, u_c \in \mathcal{U}_c} u_c(h, h') < \infty, \text{ where } D^n \text{ is the set of doctors in } \hat{\Gamma}^n.$$

The first four conditions require that the sequence of matching games induce a regular sequence of random markets. More specifically, if we define a sequence of random markets by focusing on ordinal preferences represented by utility functions in the matching games, then the induced sequence of random markets is regular in the sense of Definition 1. The last condition (5) does not have a counterpart in the definition of regular markets. It requires that utilities are bounded across matching games in the given sequence.

**Theorem 2.** *Consider a regular sequence of matching games. For any  $\epsilon > 0$ , there exists  $n$  such that truth-telling is an  $\epsilon$ -Bayes Nash equilibrium for every matching game in that sequence with more than  $n$  hospitals.*

We defer the proof to the Appendix and describe the main argument here. First consider incentives for single doctors. We begin by recalling that the doctor-proposing deferred acceptance algorithm, which is Step 1 of the sequential couples algorithm and hence part of mechanism  $\varphi$ , is strategy-proof for doctors. Next consider Steps 2 and 3 in the sequential couples algorithm, as couples are added to the market. If the market is large and doctor's preference lists have bounded lengths, then it can be shown that with high probability there are a large number of hospitals with vacant positions. In such a circumstance, at each instance of Steps 2 and 3, it is much more likely for the current applicant(s) to apply to a (pair of) vacant position(s) than to a hospital where any particular single doctor is tentatively assigned. Because there are only a finite number of applicants in Steps 2 and 3 of the algorithm, it is very unlikely for any particular single doctor to be displaced in these Steps. In other words, for each doctor, it is likely that at the end of Step 3 she receives the same matching as in the deferred acceptance algorithm. This implies that the probability that she is made strictly better off is small, and hence we conclude that a unilateral deviation cannot make her better off by a large amount.

Analysis of hospitals' incentives is more involved. The first challenge is that even the doctor-proposing deferred acceptance algorithm is not strategy-proof for hospitals, unlike for doctors. Thus the large size of the market is first exploited to establish that hospitals are unlikely to benefit by misreporting under the deferred acceptance algorithm. To see this, we first observe that a hospital has an opportunity to profitably manipulate its preferences only when a strategic rejection by a hospital causes a chain of applications and rejections that benefits the manipulating hospital. More specifically, if a hospital strategically rejects some doctors, then these doctors apply to other hospitals, displacing other doctors, who apply to yet other hospitals, and so on. This chain may cause some doctor rejected

elsewhere to apply to the original manipulating hospital, and the hospital may be made better off if this new applicant is desirable. In fact, one can show that whenever a hospital’s manipulation is successful, there is a corresponding chain of applications and rejections described above, so showing that there is no such chain suffices for showing manipulation is not profitable (Kojima and Pathak (2009) formalize and prove this claim). Now, in a large market, there is a high probability that there will be many hospitals with vacant positions at the conclusion of the algorithm. Hence the doctors who are strategically rejected (or those who are rejected by them and so on) are likely to apply to those vacant positions and be accepted rather than apply to the original manipulating hospital. As a result, the manipulating hospital is unlikely to be made better off in the deferred acceptance algorithm. Moreover, an argument similar to that for the single doctors can be made, which shows that the mechanism  $\varphi$  is unlikely to change the assignment for the hospital after the deferred acceptance algorithm if the market is sufficiently large. These arguments establish that hospitals do not benefit by a large amount by misreporting when the market size is large.

To investigate incentives for couples, note that when a couple is tentatively matched for the first time during the algorithm, they are matched to their most preferred hospital(s) among the available ones at that step. Thus any misreport of preferences results in a weakly less preferred tentative matching at that step. Moreover, if the market is sufficiently large, an argument similar to those for single doctors and hospitals establishes that the final matching for the couple is likely to be identical to the one obtained above. This implies that a couple cannot improve their expected utility by a large amount by misreporting their preferences.

## 7 Conclusion

This paper is a first step towards understanding complementarities in matching markets. We investigate this issue by studying couples in labor market clearinghouses, a phenomenon that has grown in importance as the number of dual-career households have become an important part of the labor force. Even though a stable matching does not necessarily exist in the presence of couples, as long as the complementarities caused by couples are small in an appropriate way, the market still has a stable matching with a high probability. Our second main result is that in a large game induced by a particular mechanism, there is an epsilon equilibrium at which all participants – single doctors, couples, and hospitals – submit their preferences truthfully.

One of the earlier findings on large matching markets is that when the length of preference lists of doctors is limited (e.g. by the number of interviews it is possible to attend) the number of stable matchings becomes relatively small in large markets. This might suggest that if a stable matching does

not exist in a finite market, then it is unlikely to exist in a large market. Our analysis finds, on the contrary, that largeness can be used to overcome non-existence in the finite market. Methodologically, our analysis suggests that large market arguments may be useful in generating new kinds of results in the presence of complementarities, which have traditionally proven elusive in market design problems.

It may be possible to relax some of the assumptions we have made about how markets become large (and the proportion of couples remain small). This extension may require making stronger assumptions about couples' preferences than we have made as might be justified from the empirical evidence. It is an open question whether the conclusions of this paper hold if we allow the number of couples to grow at the same order as the market size. This difficult question is left for future work.<sup>27</sup>

We have complemented our theoretical results with analysis of data from the market for clinical psychologists. The data suggested empirically relevant modeling assumptions. In every year of the data it is possible to find a stable matching with respect to the stated preferences using the mechanism we analyze. Given our result that truth-telling is an approximate best response in large markets, this empirical result may suggest that there exists a stable matching with respect to the true preferences of participants. Since the mechanism we analyze is similar to the actual procedures used in markets such as the NRMP for American medical residents, our results help explain why some mechanisms in practice provide a stable matching with high probability even in the presence of couples, at least in large markets without too many couples.

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<sup>27</sup>As we have highlighted, the conclusions hold as long as the number of couples grow reasonably slowly, specifically more slowly than the square root of the number of hospitals. Tight results on the rates of convergence are often elusive in papers on large markets, although there are notable advancements such as Rustichini et al. (1994) in a different context, namely in double auction markets.

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## A Appendix

### A.1 Formal Definitions

#### A.1.1 Definition of Responsive Preferences

Let  $h \in H$  and  $\kappa_h$  be a positive integer. We say that preference relation  $\succeq_h$  is **responsive with capacity**  $\kappa_h$  if

- (1) For any  $D' \subseteq D$  with  $|D'| \leq \kappa_h$ ,  $d \in D \setminus D'$  and  $d' \in D'$ ,  $D' \cup d \setminus d' \succeq_h D'$  if and only if  $d \succeq_h d'$ ,
- (2) For any  $D' \subseteq D$  with  $|D'| \leq \kappa_h$  and  $d' \in D'$ ,  $D' \succeq_h D' \setminus d'$  if and only if  $d' \succeq_h \emptyset$ , and
- (3)  $\emptyset \succ_h D'$  for any  $D' \subseteq D$  with  $|D'| > \kappa_h$ .

#### A.1.2 Definition of Individual Rationality

A matching  $\mu$  is **individually rational** if

- (1)  $\mu(s)R_s\emptyset$  for every  $s \in S$ ,
- (2)  $\mu(c)R_c(\emptyset, \emptyset)$  for every  $c \in C$ , and
- (3)  $Ch_h(\mu(h)) = \mu(h)$  for every  $h \in H$ .<sup>28</sup>

### A.2 An Alternative Definition of Stability

We offer an alternative definition of stability, which we call strong stability, than the one presented in the main text. This alternative definition allows for larger coalitions to block a matching. A matching which is strongly stable is also stable according to the definition in the main text. In the proof of Theorem 1, we establish a more general result for strong stability, and this implies existence of a stable matching as defined in the main text.

In the definition of strong stability, we consider two cases of a block as follows:

- (1) A couple-hospital pair  $(c, h) \in C \times H$  is a **block** of  $\mu$  if
  - (a)  $(h, h)P_c\mu(c)$ ,
  - (b)  $f, m \in Ch_h(\mu(h) \cup c)$  where  $c = (f, m)$ .

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<sup>28</sup>When there is a couple  $(f, m)$  with  $\{f, m\} \subseteq \mu(h)$ , we adopt a notational convention that  $Ch_h(\mu(h))$  means  $Ch_h(\mu(h) \cup (f, m) \setminus \{f, m\})$ , that is, we let hospital  $h$  to consider the existing couple as a whole when choosing the most preferred subset of doctors. Similar conventions will be used elsewhere when the choice involves a couple who are matched as a whole at the given matching.

- (2) A group of doctors  $D'$  and hospital  $h$  is a **block** of  $\mu$  if
- (a) there is no couple  $(f, m)$  such that  $\{f, m\} \subseteq D'$ ,
  - (b)  $D' \subseteq Ch_h(\mu(h) \cup D')$ ,<sup>29</sup>
  - (c) for all  $s \in D' \cap S$ , we have  $hP_s\mu(s)$ ,
  - (d)
    - i. for all  $f \in D' \cap F$  where  $c = (f, m) \in C$ ,  $(h, h')P_c\mu(c)$  for some  $h'$  and  $m \in Ch_{h'}(\mu(h') \cup m)$ ,
    - ii. for all  $m \in D' \cap M$  where  $c = (f, m) \in C$ , we have  $(h', h)P_c\mu(c)$  for some  $h'$  and  $f \in Ch_{h'}(\mu(h') \cup f)$ .

A matching is **strongly stable** if it is individually rational and there is no block as defined by conditions (1) and (2) above.

This definition allows for a couple assigned to a hospital to be blocked by two doctors (who are either single or are a member of a couple). If one of the blocking doctors is a couple member, we require that the member's partner is chosen by another hospital over its assignment and that the couple together prefer this assignment to their current assignment.

The motivation for this definition is to allow certain joint deviations to happen, but rule out more complicated deviations involving larger groups. When a couple member is part of a blocking coalition of doctors  $D'$  and hospital  $h$ , our interpretation is that hospital  $h$  is the “initiating” blocker, and any hospital  $h'$  involved to satiate the other member of a couple as the passive blocker. The reason we consider this definition is to stay close to pairwise stability, but still accommodate this particular type of blocking pair is that we think that blocking coalitions of larger size are less likely to form due to coordination issues among members. This definition also keeps the notation less burdensome, and but our main existence result continues to hold when we allow larger sets of blocking coalitions to form or employ the core as our solution concept.

Since the definition of strong stability allows for coalitions of doctors who are single or couple members to be part of blocking pairs, a strongly stable matching is stable, but not vice versa. Moreover, if each hospital has one position, strong stability is equivalent to unit-capacity stability.

Finally, strong stability is equivalent to the standard definition of (pairwise) stability when there is no couple. To see this last point, first observe that condition (1) in the definition of strong stability is irrelevant if there is no couple, as are conditions (2a) and (2d). The remaining conditions (2b) and (2c)

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<sup>29</sup>When there is a couple  $(f, m)$  with  $\{f, m\} \subseteq \mu(h)$ , then we adopt a notational convention that  $Ch_h(\mu(h) \cup D')$  means  $Ch_h(\mu(h) \cup (f, m) \setminus \{f, m\} \cup D')$ , that is, we let hospital  $h$  to consider the existing couple as a whole when choosing the most preferred subset of doctors. Similar convention will be used elsewhere when the choice involves a couple who are matched as a whole at the given matching.

are equivalent to the nonexistence of a blocking pair under the assumption that hospital preferences are responsive. Thus this, together with individual rationality, is equivalent to the standard pairwise stability concept.

### A.3 Proof of Theorem 1

Let  $(H, S, C, (\succeq_h)_{h \in H}, (R_i)_{i \in S \cup C})$  be a matching market.

#### Step 1: Doctor-Proposing Deferred Acceptance Algorithm

Apply the doctor-proposing deferred acceptance algorithm to the sub-market without couples:  $(H, S, (\succeq_h)_{h \in H}, (R_s)_{s \in S})$ .

#### Step 2: Sequential Couples Algorithm

##### Algorithm 2. SEQUENTIAL COUPLES ALGORITHM

(1) Initialization:

Let matching  $\mu$  be the output of the deferred acceptance algorithm in the sub-market without couples.

(2) Iterate through couples: set  $C^0 = C$ ,  $i = 0$  and  $B = \emptyset$ .

- (a) If  $C^i$  is empty, then go to Step 3. Otherwise, pick some couple  $c = (f, m) \in C^i$ . Let  $C^{i+1} = C^i \setminus c$  and increment  $i$  by one.
- (b) Let couple  $c$  apply to their most preferred pair of hospitals  $(h, h') \in H \times H$  that has not rejected them yet.
  - i. If such a hospital (pair) does not exist, modify matching  $\mu$  such that couple  $c$  is unassigned and then go to Step 2a.
  - ii. If such a hospital (pair) exists, then if either hospital  $h$  or hospital  $h'$  has previously been applied to by a member (or both members) of any couple different from  $c$ , then terminate the algorithm.
  - iii. Otherwise,
    - A. If  $h = h' \neq \emptyset$  and  $\{f, m\} \subseteq Ch_h(\mu(h) \cup c)$ , then modify matching  $\mu$  by assigning  $(f, m)$  to hospital  $h$  and having  $h$  reject

$$(\mu(h) \cup f \cup m) \setminus Ch_h(\mu(h) \cup c).$$

Add the rejected single doctors (if any) to  $B$  and go to Step 2a.

- B. If  $h \neq h'$ ,  $f \in Ch_h(\mu(h) \cup f)$ , and  $m \in Ch_{h'}(\mu(h') \cup m)$ , then modify matching  $\mu$  by assigning  $f$  to  $h$  and  $m$  to  $h'$ , having hospital  $h$  reject

$$(\mu(h) \cup f) \setminus Ch_h(\mu(h) \cup f),$$

and having hospital  $h'$  reject

$$(\mu(h') \cup m) \setminus Ch_{h'}(\mu(h') \cup m).$$

Add the rejected single doctors (if any) to  $B$  and go to Step 2a.

- C. Otherwise, let hospital  $h$  and hospital  $h'$  reject the application by couple  $c$  and go to Step 2b.

- (3) *Iterate through rejected single doctors:* set  $B^1 = B$  and  $j = 1$ .

Round  $j$ :

- (a) If  $B^j$  is empty, then terminate the algorithm.  
(b) Otherwise, pick some single doctor  $s$  in  $B^j$ . Let  $B^{j+1} = B^j \setminus s$  and increment  $j$  by one.

*Iterate through the rank order lists of single doctors:*

- i. If single doctor  $s$  has applied to every acceptable hospital, then modify matching  $\mu$  such that  $s$  is unassigned and go to Step 3a.
- ii. If not, then let  $\hat{h}$  be the most preferred hospital ranked by single doctor  $s$  among those which  $s$  has not yet applied to previously (either in the doctor-proposing deferred acceptance algorithm or within this algorithm.)
- iii. If there is no couple member who has ever applied to hospital  $\hat{h}$ , then there are three cases:
  - A. If hospital  $\hat{h}$  has a vacant position and  $s$  is acceptable to  $\hat{h}$ , then modify matching  $\mu$  such that single doctor  $s$  is assigned to  $\hat{h}$  and go to Step 3a.
  - B. If either hospital  $\hat{h}$  prefers each of its current mates to single doctor  $s$  and there is no vacant position or  $s$  is unacceptable to  $\hat{h}$ , then  $\hat{h}$  rejects  $s$  and go to Step 3(b)i.
  - C. If hospital  $\hat{h}$  prefers single doctor  $s$  to one of its current mates and there is no vacant position, then modify matching  $\mu$  such that  $s$  is assigned to  $\hat{h}$ . Hospital  $\hat{h}$

rejects the least preferred doctor currently assigned there:

$$(\mu(\hat{h}) \cup s) \setminus Ch_{\hat{h}}(\mu(\hat{h}) \cup s).$$

With abuse of notation, denote this rejected doctor  $s$  and go to Step 3(b)i.

- iv. If there is a couple member who has ever applied to hospital  $\hat{h}$  previously within this algorithm, then terminate the algorithm.

The sequential couples algorithm terminates at Step 2(b)ii (when a couple member proposes to a hospital which has already been proposed to by another couple), Step 3a (when all couples and single doctors are assigned), or Step 3(b)iv (when a single doctor proposes to a hospital which was previously applied to by a couple member). We say that the algorithm **succeeds** if it terminates at Step 3a.

**Lemma 1.** *If the sequential couples algorithm succeeds, then the resulting matching is stable.*

The proof of this lemma is similar to the proof of the existence of a stable matching by Gale and Shapley (1962) in the college admissions model. The main difference is that when the sequential couples algorithm succeeds, we must verify that there are no blocking pairs including pairs which may involve members of a couple.

*Proof of Lemma 1.* We prove that the matching that results when the sequential couples algorithm succeeds is strongly stable (defined in Section A.2). Establishing this fact implies that the matching is stable since a strongly stable matching is a stable matching.<sup>30</sup>

Suppose that Algorithm 2 succeeds, producing matching  $\mu$ . First,  $\mu$  is individually rational since all doctors who are single or couple members have applied only to acceptable hospitals (hospital pairs for couples), and hospitals have accepted only acceptable doctors only up to their capacities in each step of Algorithms 1 and 2.

Next, to show that there is no block of matching  $\mu$ , fix a hospital  $h \in H \setminus \emptyset$ .

- (1) Suppose that there exists no couple  $(f, m) \in C$  such that  $\{f, m\} \subseteq \mu(h)$ .
  - (a) Assume, for contradiction, that there exists a set of doctors  $D' \subseteq D$  such that hospital  $h$  and  $D'$  block  $\mu$ , where there is no couple  $(f, m) \in C$  such that  $\{f, m\} \subseteq D'$ . Since doctor  $d$  is part of a block,

$$dP_h \emptyset \quad \text{for every } d \in D'.$$

There are two cases to consider depending on whether  $D'$  contains any single doctors.

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<sup>30</sup>Since the rest of the analysis builds on this lemma, this stronger result allows us to extend our main results when we replace stability with strong stability as the solution concept.

- i. Suppose that there is a single doctor in  $D'$ . Then each single doctor  $s \in D' \cap S$  is rejected by hospital  $h$  at some point of either Algorithm 1 or 2 since  $hP_s\mu(s)$ . The tentative assignment of hospital  $h$  at a step when single doctor  $s$  is rejected, denoted  $\tilde{\mu}(h)$ , satisfies

$$|\tilde{\mu}(h)| = \kappa_h \quad \text{and} \quad dP_h s \text{ for all } d \in \tilde{\mu}(h),$$

because  $sP_h\emptyset$ . Since, at each of later steps of both Algorithms, hospital  $h$  replaces a tentatively matched doctor only when a more preferred doctor applies, it follows that

$$|\mu(h)| = \kappa_h \quad \text{and} \quad dP_h s \text{ for all } d \in \mu(h).$$

This contradicts the assumption that hospital  $h$  and  $D'$  block matching  $\mu$ .

- ii. Suppose there are no single doctors in  $D'$ . Then there exists a member of some couple in  $D'$ . Without loss of generality, assume that there is some  $f \in D'$  where  $c = (f, m) \in C$ . Since  $(h, h')P_c\mu(c)$  for some  $h' \in H$ , couple  $c$  was rejected by the hospital pair  $(h, h')$  at some point of Algorithm 2. Let  $\tilde{\mu}(h)$  and  $\tilde{\mu}(h')$  be the tentative assignments for hospital  $h$  and hospital  $h'$  at that step, respectively. Because couple  $c$  was rejected at this step, it follows that either

$$|\tilde{\mu}(h)| = \kappa_h \quad \text{and} \quad dP_h f \text{ for all } d \in \tilde{\mu}(h),$$

or  $h' \neq \emptyset$  and we have that

$$|\tilde{\mu}(h')| = \kappa_{h'} \quad \text{and} \quad dP_{h'} m \text{ for all } d \in \tilde{\mu}(h').$$

Since, at each of later steps, both hospital  $h$  and hospital  $h'$  (if  $h' \neq \emptyset$ ) replace a tentatively matched doctor only when a more preferred doctor applies, it follows that either

$$|\mu(h)| = \kappa_h \quad \text{and} \quad dP_h f \text{ for all } d \in \mu(h),$$

or  $h' \neq \emptyset$  and we have that

$$|\mu(h')| = \kappa_{h'} \quad \text{and} \quad dP_{h'} m \text{ for all } d \in \mu(h').$$

This contradicts the assumption that  $f \in D'$  and  $D'$  block matching  $\mu$  with hospital  $h$ .

- (b) Consider a couple  $c = (f, m)$  such that  $(h, h)P_c\mu(c)$ . By definition of Algorithm 2, the couple was rejected by the hospital pair  $(h, h)$  at some point in the Algorithm. Denote the

matching at that point by  $\tilde{\mu}$ . It follows that

$$Ch_h(\tilde{\mu}(h) \cup c) = \tilde{\mu}(h) \quad \text{and} \quad f, m \notin \tilde{\mu}(h).$$

Since the sequential couples algorithm succeeds, no other doctor applies to hospital  $h$  after the step where couple  $c$  is rejected by  $(h, h)$ . As a result,

$$\mu(h) = \tilde{\mu}(h).$$

Therefore,

$$f, m \notin \mu(h) \quad \text{and} \quad \mu(h) = Ch_h(\mu(h) \cup c),$$

which contradicts the assumption that couple  $c$  and hospital  $h$  block matching  $\mu$ .

- (2) Suppose that there exists a couple  $(f, m) \in C$  such that  $\{f, m\} \subseteq \mu(h)$  and there is a block of matching  $\mu$  involving hospital  $h$ . The assumption that the sequential couples algorithm succeeds implies that there is no couple  $c \neq (f, m)$  and  $h' \in H$  such that  $(h, h')P_c\mu(c)$  or  $(h', h)P_c\mu(c)$ . This is because the algorithm terminates in Step 3 if two or more distinct couple members apply to the same hospital during the algorithm. Thus, the set of doctors  $D'$  that blocks matching  $\mu$  with hospital  $h$  is composed solely of single doctors. This means that

$$sP_h\emptyset \quad \text{for every } s \in D', \tag{1}$$

$$hP_s\mu(s) \quad \text{for every } s \in D', \tag{2}$$

for otherwise a single doctor  $s$  is not part of a block. Let  $\tilde{\mu}$  be the matching that is the result of the doctor-proposing deferred acceptance algorithm in the sub-market excluding couples. Then

$$hR_s\tilde{\mu}(s) \quad \text{for every } s \in D', \tag{3}$$

because otherwise, in light of (2), single doctor  $s$  will have applied to hospital  $h$  in Step 3 of the sequential couples algorithm, causing the algorithm to fail. Moreover,

$$\mu(h) \succ_h \tilde{\mu}(h), \tag{4}$$

because otherwise hospital  $h$  would not have accepted new applicants in Step 2 of the sequential

couples algorithm, resulting in matching with  $\mu(h)$ . Furthermore,

$$dP_h s \quad \text{for every } d \in \tilde{\mu}(h) \cap \mu(h) \text{ and } s \in D', \quad (5)$$

because

- (a) for any  $s \in D' \cap \tilde{\mu}(h)$ , single doctor  $s$  was rejected in Step 2 of the sequential couples algorithm at the instance when the couple  $(f, m)$  applied to hospital  $h$ . We now show  $dP_h s$ . Suppose, to the contrary, that  $sR_h d$  for some  $d \in \tilde{\mu}(h) \cap \mu(h)$ . Then  $sP_h d$  because  $s \neq d$  and preferences are strict. This relation and responsiveness of  $\succeq_h$  imply

$$\mu(h) \cup s \setminus d \succ_h \mu(h).$$

Moreover,

$$\mu(h) \cup s \setminus d \in \mathcal{A}(\tilde{\mu}(h) \cup (f, m)).$$

These facts contradict

$$\mu(h) = Ch_h(\tilde{\mu}(h) \cup (f, m)),$$

which follows from the definition of the sequential couples algorithm. Hence,

$$dP_h s \quad \text{for all } d \in \tilde{\mu}(h) \cap \mu(h).$$

- (b) for any  $s \in D' \setminus \tilde{\mu}(h)$ , relation (3) implies  $hP_s \tilde{\mu}(s)$ , so stability of matching  $\tilde{\mu}$  in the sub-market without couples (which coincides with both our stability definition in Section 3.1 and strong stability definition in Appendix A.2) implies

$$dP_h s \quad \text{for all } d \in \tilde{\mu}(h).$$

Let  $\mu'(h)$  be the assignment for hospital  $h$  when  $D'$  and hospital  $h$  block matching  $\mu$ . That is,

$$\mu'(h) = Ch_h(\mu(h) \cup D'). \quad (6)$$

Relation (6) and the definition of  $Ch_h(\cdot)$  imply

$$\mu'(h) \succ_h \mu(h). \quad (7)$$

Relations (1) and (2) imply that

$$|\mu(h)| = \kappa_h.$$

Therefore, to block matching  $\mu$  with  $D'$ , hospital  $h$  should reject some doctors in  $\mu(h)$ . If any doctor  $d \in \tilde{\mu}(h) \cap \mu(h)$  is rejected while some  $s \in D'$  is accepted to produce  $\mu'(h)$ , then

$$\mu'(h) \cup d \setminus s \succ_h \mu'(h)$$

by responsiveness of  $\succeq_h$  and relation (5), but this contradicts (6). Hence, relation (5) implies that it should be exactly couple  $(f, m)$  that is rejected by hospital  $h$  when hospital  $h$  and  $D'$  block  $\mu(h)$ . Since  $(f, m)$  is the only couple in  $\mu(h)$  and it is not in  $\mu'(h)$ ,

$$\mu'(h) \subseteq \tilde{\mu}(h) \cup D' \subseteq S.$$

Since  $hR_s \tilde{\mu}(s)$  for every single doctor  $s \in \mu'(h)$  by relation (3) and  $\mu'(h) \subseteq S$ , it follows that

$$\tilde{\mu}(h) \succeq_h \mu'(h), \tag{8}$$

because otherwise matching  $\tilde{\mu}$  would be unstable in the sub-market without couples. Applying relations (4), (8), and then (7), we obtain

$$\mu(h) \succ_h \tilde{\mu}(h) \succeq_h \mu'(h) \succ_h \mu(h),$$

a contradiction. □

The rest of our argument uses Lemma 1 to compute how often Algorithm 2 succeeds when singles and couples draw their preferences according to the processes described in Section 4.1.

In the next two steps of the proof, we define versions of the deferred acceptance algorithm and the sequential couples algorithm where single doctors draw their preferences iteratively within the steps of the algorithms. This representation of the two algorithms proves useful for our analysis.

### Step 3: Define Stochastic Deferred Acceptance Algorithm

#### Algorithm 3. STOCHASTIC DOCTOR-PROPOSING DEFERRED ACCEPTANCE ALGORITHM

- (1) Initialization: Let  $l = 1$ . For every  $s \in S$ , let  $A_s = \emptyset$  and order the single doctors in an arbitrarily fixed manner.

- (2) Choosing the applicant:
  - (a) If  $l \leq |S|$ , then let  $s$  be the  $l^{\text{th}}$  single doctor and increment  $l$  by one.
  - (b) If not, then terminate the algorithm.
- (3) Choosing the applied:
  - (a) If  $|A_s| \geq k$ , then return to Step 2.
  - (b) If not, select hospital  $h$  randomly from distribution  $\mathcal{P}^n$  until  $h \notin A_s$ , and add  $h$  to  $A_s$ .
- (4) Acceptance and/or rejection:
  - (a) If hospital  $h$  prefers each of its current mates to single doctor  $s$  and there is no vacant position, then hospital  $h$  rejects single doctor  $s$ . Go to Step 3.
  - (b) If hospital  $h$  has a vacant position or it prefers single doctor  $s$  to one of its current mates, then hospital  $h$  accepts single doctor  $s$ . Now if hospital  $h$  had no vacant position before accepting single doctor  $s$ , then hospital  $h$  rejects the least preferred doctor among those who were matched to hospital  $h$ . Let this doctor be  $s$  and go to Step 3. If hospital  $h$  had a vacant position, then go back to Step 2.

$A_s$  records hospitals that single doctor  $s$  has already drawn from  $\mathcal{P}^n$ . When  $|A_s| = k$  is reached,  $A_s$  is the set of hospitals acceptable to single doctor  $s$ .

Let  $\mu$  be the matching that is produced when Algorithm 3 terminates. Under the doctor proposing deferred acceptance algorithm, a single doctor's application to her  $t^{\text{th}}$  most preferred hospital is independent of her preferences after  $(t + 1)^{\text{th}}$  choice on. Therefore matching  $\mu$  is stable for the market consisting of single doctors, any of their realized preference profiles which could follow from completing the draws for random preferences, the hospitals and their (arbitrarily fixed) preferences.

#### Step 4: Define Stochastic Sequential Couples Algorithm

Suppose that at the conclusion of Algorithm 3, we obtain matching  $\mu$ . The **stochastic sequential couples algorithm** is a version of Algorithm 2 where single doctor preferences are drawn iteratively, and is defined as follows:

#### Algorithm 4. STOCHASTIC SEQUENTIAL COUPLES ALGORITHM

- (1) Initialization:

- (a) Keep all preference lists generated in Algorithm 3. Also, for each single doctor  $s \in S$ , let  $A_s$  be the set generated at the end of Algorithm 3. Let the matching  $\mu$  be the initial matching of the algorithm.
- (b) For each couple  $c = (f, m) \in C$ , construct the couples' preferences  $P_c$  according to the process defined in Section 4.1.
- (2) *Iterate through couples*, set  $C^0 = C$ ,  $i = 0$ , and  $B = \emptyset$ .
- (a) If  $C^i$  is empty, then go to Step 3. Otherwise, pick some couple  $c = (f, m) \in C^i$ . Let  $C^{i+1} = C^i \setminus c$  and increment  $i$  by one.
- (b) Let couple  $c$  apply to their most preferred pair  $(h, h') \in H \times H$  that has not rejected them yet.
- i. If such a hospital (pair) does not exist, modify matching  $\mu$  such that couple  $c$  is unassigned and then go to Step 2a.
  - ii. If such a hospital (pair) exists, then if either hospital  $h$  or hospital  $h'$  has previously been applied to by a member (or both members) of any couple different from  $c$ , then terminate the algorithm.
  - iii. Otherwise,
    - A. If  $h = h' \neq \emptyset$  and  $\{f, m\} \subseteq Ch_h(\mu(h) \cup c)$ , then modify matching  $\mu$  by assigning  $(f, m)$  to hospital  $h$  and having  $h$  reject
 
$$(\mu(h) \cup f \cup m) \setminus Ch_h(\mu(h) \cup c).$$

Add the rejected single doctors (if any) to  $B$  and go to Step 2a.
    - B. If  $h \neq h'$ ,  $f \in Ch_h(\mu(h) \cup f)$ , and  $m \in Ch_{h'}(\mu(h') \cup m)$ , then modify matching  $\mu$  by assigning  $f$  to  $h$  and  $m$  to  $h'$ , having hospital  $h$  reject
 
$$(\mu(h) \cup f) \setminus Ch_h(\mu(h) \cup f),$$

and having hospital  $h'$  reject

$$(\mu(h') \cup m) \setminus Ch_{h'}(\mu(h') \cup m).$$

Add the rejected single doctors (if any) to  $B$  and go to Step 2a.
    - C. Otherwise, let hospital  $h$  and hospital  $h'$  reject the application by couple  $c$  and go to Step 2b.

(3) Iterate through rejected single doctors, set  $B^1 = B$  and  $j = 1$ .

Round  $j$ :

- (a) If  $B^j$  is empty, then terminate the algorithm.
- (b) Otherwise, pick some single doctor  $s$  in  $B^j$ . Set  $B^{j+1} = B^j \setminus s$  and increment  $j$  by one.

*Iterate through the single doctor's rank order list (call this iteration "Round  $j$ ")*

- i. If  $|A_s| \geq k$ , then go to Step 3a.
- ii. If not, select hospital  $\hat{h}$  randomly from distribution  $\mathcal{P}^n$  until  $\hat{h} \notin A_s$ , and add  $\hat{h}$  to  $A_s$ .
- iii. If there is no couple member who has ever applied to hospital  $\hat{h}$ , then there are three cases:
  - A. If hospital  $\hat{h}$  has a vacant position, then modify matching  $\mu$  such that single doctor  $s$  is assigned to  $\hat{h}$  and go to Step 3a.
  - B. If either hospital  $\hat{h}$  prefers each of its current mates to single doctor  $s$  and there is no vacant position or  $s$  is unacceptable to  $\hat{h}$ , then  $\hat{h}$  rejects  $s$  and go to Step 3(b)i.
  - C. If hospital  $\hat{h}$  prefers single doctor  $s$  to one of its current mates and there is no vacant position, then modify matching  $\mu$  such that  $s$  is assigned to  $\hat{h}$ . Hospital  $\hat{h}$  rejects the least preferred doctor currently assigned there

$$(\mu(\hat{h}) \cup s) \setminus Ch_{\hat{h}}(\mu(\hat{h}) \cup s).$$

With abuse of notation, denote this rejected doctor  $s$  and iterate through her rank order list by going to Step 3(b)i.

- iv. If there is a couple member who has ever applied to hospital  $\hat{h}$  before, then terminate the algorithm.

The algorithm above terminates at Step 2(b)ii or Step 3a or Step 3(b)iv. Similarly to Algorithm 2, we say that Algorithm 4 **succeeds** if it terminates at Step 3a.

To establish Theorem 1, we investigate how often the algorithm succeeds, as single doctors draw hospitals from distribution  $\mathcal{P}^n$  and couples draw hospitals from  $\mathcal{Q}^n$ . First observe for any random market in a regular sequence,

$$\Pr[\text{Algorithm 2 succeeds}] = \Pr[\text{Algorithm 4 succeeds}].$$

That is, the probability of the algorithm's success is identical whether random preferences are drawn at once in the beginning or they are drawn one at a time during the execution of the algorithm. The latter expression is useful since we can investigate the procedure step by step, utilizing conditional probabilities and conditional expectations. Thus we focus on the behavior of Algorithm 4 as the market size grows in the remainder of the proof.

Let  $Y_n$  be a random variable which counts the number of hospitals that are listed on no single doctor preference list at the end of Algorithm 3.<sup>31</sup> The next step of the argument provides a lower bound on  $Y_n$  at the conclusion of Algorithm 3.

### Step 5: A large number of hospitals have vacancies

**Lemma 2.** *For any  $n > 4k$ ,*

$$E[Y_n] \geq \frac{n}{2} e^{-16\lambda k}.$$

*Proof.* We order the hospitals in terms of decreasing popularity with respect to  $\mathcal{P}^n$ : label  $H = \{h_1, h_2, \dots, h_n\}$  with  $p_{h_\ell} \geq p_{h_{\ell+1}}$  for all  $\ell \in \{1, \dots, n-1\}$ . If hospitals have equal popularity, we order them in some fixed manner. Let the probability mass for the first  $k$  hospitals be:

$$\pi = \sum_{\ell=1}^k p_{h_\ell}.$$

Fix a single doctor and denote her  $i^{\text{th}}$  most preferred hospital by  $h_{(i)}$ , if it has been drawn at the conclusion of Algorithm 3. For any  $i \leq k$ , the definition of  $\pi$  implies

$$\sum_{j=1}^{i-1} p_{h_{(j)}} \leq \pi.$$

Thus for any  $i \leq k$ , the conditional probability that  $h$  is not the single doctor's  $i^{\text{th}}$  choice given the events that her first  $(i-1)$  choices are  $h_{(1)}, \dots, h_{(i-1)}$ , her  $i^{\text{th}}$  choice is drawn, and  $h_{(j)} \neq h$  for all  $j \leq i-1$ , is bounded from below by

$$1 - \frac{p_h}{1 - \sum_{j=1}^{i-1} p_{h_{(j)}}} \geq 1 - \frac{p_h}{1 - \pi}. \quad (9)$$

Let  $E_h$  be the event that  $h \notin A_s$  for every  $s \in S$  at the end of Algorithm 3. Since at most  $\lambda nk$  draws

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<sup>31</sup>We abuse notation and denote a random variable and its realization by the same letter when there is no confusion.

are made in total by all single doctors from  $\mathcal{P}^n$  in Algorithm 3, inequality (9) implies that

$$\Pr(E_h) \geq \left(1 - \frac{p_h}{1 - \pi}\right)^{\lambda nk}. \quad (10)$$

If hospital index  $\ell > k$ , there are at least  $\ell - k$  hospitals in  $H \setminus \{h_1, \dots, h_k\}$  that are at least as popular as  $h_\ell$  according to  $\mathcal{P}^n$ , so we obtain

$$\begin{aligned} \pi + (\ell - k)p_{h_\ell} &\leq 1 \\ \iff p_{h_\ell} &\leq \frac{1 - \pi}{\ell - k}. \end{aligned} \quad (11)$$

The last two inequalities, (10) and (11), imply

$$\Pr(E_{h_\ell}) \geq \left(1 - \frac{1}{\ell - k}\right)^{\lambda nk}. \quad (12)$$

We next show that for any  $\ell > 2k$ ,

$$\left(1 - \frac{1}{\ell - k}\right)^{\lambda nk} \geq e^{-2\lambda nk/(\ell - k)}. \quad (13)$$

To see that (13) holds, first note that

$$\left(1 - \frac{1}{\ell - k}\right)^{\lambda nk} \geq e^{-2\lambda nk/(\ell - k)} \iff 1 - \frac{1}{\ell - k} - e^{-2/(\ell - k)} \geq 0.$$

Now, define a function  $g(x) = 1 - x - e^{-2x}$ . This function  $g$  is concave, and  $g(0) = 0$  and  $g(1/2) = 1/2 - 1/e > 0$ . Therefore  $g(x) \geq 0$  for any  $x \in [0, 1/2]$ . Since  $\ell > 2k$  and  $k$  is a positive integer, we have  $\ell - k > k \geq 1$ . Since  $\ell - k$  is an integer, we obtain  $\ell - k \geq 2$  and hence  $1/(\ell - k) \in [0, 1/2]$ . Therefore

$$1 - 1/(\ell - k) - e^{-2/(\ell - k)} = g(1/(\ell - k)) \geq 0,$$

establishing inequality (13).

Moreover, for any  $\ell > 2k$ ,

$$e^{-2\lambda nk/(\ell - k)} \geq e^{-4\lambda nk/\ell}. \quad (14)$$

Combining inequalities (12), (13), and (14), we obtain:

$$\Pr(E_{h_\ell}) \geq e^{-4\lambda nk/\ell}.$$

Using this inequality, for any  $n > 4k$ , we have

$$E[Y_n] = \sum_{\ell=1}^n \Pr(E_{h_\ell}) \geq \sum_{\ell=2k}^n e^{-4\lambda nk/\ell} \geq \sum_{\ell=\lfloor n/2 \rfloor + 1}^n e^{-8\lambda nk/\ell} \geq \frac{n}{2} e^{-16\lambda k},$$

where  $\lfloor x \rfloor$  is the largest integer not greater than  $x$ .

□

**Step 6: Algorithm 4 succeeds with high probability.**

Let  $\bar{C} = bn^a$  denote the upper bound on the number of couples in the random market  $\tilde{\Gamma}^n$ .

**Lemma 3.** *For any sufficiently large  $n$  and any matching  $\mu$ ,*

$$\begin{aligned} & \Pr \left[ \text{Algorithm 4 succeeds} \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu \right] \\ & \geq \left( 1 - \frac{2k\bar{C}r}{n} \right)^{2k\bar{C}} \cdot \left( 1 - \frac{8rk\bar{C}}{E[Y_n]} \right)^{2\bar{C}}, \end{aligned}$$

*if the conditioning event has a strictly positive probability.*

*Proof.* First, consider the event that Algorithm 4 does not terminate at Step 2(b)ii so that the algorithm reaches Step 3. For that event to happen it is enough for the following event to happen: for any two doctors  $d, d' \in F \cup M$  with  $d \neq d'$ , there is no hospital  $h \in H$  that is listed by both  $d$  and  $d'$  as an acceptable hospital. This is sufficient because our assumption on  $\rho$  implies that at most one couple member will apply to  $h$ .

Suppose  $\{d_1, \dots, d_{\ell-1}\} \in F \cup M$  are such that there exists no  $h \in H$  listed by any pair of doctors in  $\{d_1, \dots, d_{\ell-1}\} \in F \cup M$ . Furthermore, fix a doctor  $d_\ell \in F \cup M \setminus \{d_1, \dots, d_{\ell-1}\}$  and assume that her first  $i-1$  choices  $\{h_{(1)}, h_{(2)}, \dots, h_{(i-1)}\}$  have no intersection with hospitals listed by the set of doctors  $\{d_1, \dots, d_{\ell-1}\}$ . The conditional probability that her  $i^{\text{th}}$  choice  $h_{(i)}$  does not have an overlap with any of the previously picked hospital is at least

$$1 - \sum_{h: h \text{ is listed by some doctor in } \{d_1, \dots, d_{\ell-1}\}} q_h - \sum_{j=1}^{i-1} q_{h_{(j)}}. \quad (15)$$

Condition (4) of Definition 1 implies that

$$q_h \leq r q_{h'} \quad \text{for all } h, h' \in H.$$

Adding these inequalities across hospitals  $h' \in H$ , we have

$$n q_h \leq r \sum_{h' \in H} q_{h'} \quad \text{for each } h.$$

As a result,

$$q_h \leq \frac{r \sum_{h' \in H} q_{h'}}{n} \leq \frac{r}{n}.$$

Since there are at most  $\bar{C}$  couples and each member of a couple lists at most  $k$  distinct hospitals, expression (15) is bounded from below by

$$1 - \frac{2k\bar{C}r}{n}. \quad (16)$$

Recall that there are at most  $\bar{C}$  couples and each member of the couple lists at most  $k$  distinct hospitals. Expression (16) implies that the probability that for any  $d, d' \in F \cup M$  with  $d \neq d'$ , there is no hospital  $h \in H$  that is listed by both  $d$  and  $d'$  as one of their acceptable hospitals is at least

$$\left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}}, \quad (17)$$

which is positive for  $n$  sufficiently large. Expression (17) provides a lower bound of the probability that the algorithm does not terminate at Step 2(b)ii so that the algorithm reaches Step 3.

Next, we consider what happens in Step 3 assigning single doctors in the set  $B$ , conditional on the same events assumed so far and in addition that all couples are tentatively matched without the algorithm being terminated at Step 2(b)ii.

Condition (4) of Definition 1 implies that

$$p_{h'} \geq p_h/r \quad \text{for any } h, h' \in H.$$

Also observe that there are at most  $2k\bar{C}$  hospitals that are listed by a couple member in  $F \cup M$ . Denote this set of hospitals by  $H_1$  and note that

$$\sum_{h \in H_1} q_h \leq 2k\bar{C}r \cdot \min_{h \in H} \{q_h\}.$$

Moreover, there are at least  $Y_n - 2k\bar{C}$  hospitals (which is positive if  $n$  is sufficiently large and  $Y_n > \frac{E[Y_n]}{2}$ ) with vacant positions and not listed by any couple member at the beginning of Step 3 (since there are at least  $Y_n$  hospitals with vacant positions at the beginning of Step 2 and at most  $2\bar{C}$  hospitals are listed by couple members). Denote this set of hospitals by  $H_2$  and note that

$$\sum_{h \in H_2} q_h \geq (Y_n - 2k\bar{C}) \cdot \min_{h \in H} \{q_h\}.$$

We are interested in computing the probability that Round 1 of Step 3 ends at 3(b)iiiA as a single doctor applies to some hospital with vacant positions not listed by any couple member (rather than applying to a hospital that is listed by a couple member). This probability is bounded below by:

$$1 - \frac{\sum_{h \in H_1} q_h}{\sum_{h \in H_1} q_h + \sum_{h \in H_2} q_h} \geq 1 - \frac{2k\bar{C}}{\frac{Y_n - 2k\bar{C}}{r} + 2k\bar{C}} > 1 - \frac{2k\bar{C}}{\frac{E[Y_n]/2 - 2k\bar{C}}{r} + 2k\bar{C}}. \quad (18)$$

Now assume that all Rounds  $1, \dots, j-1$  end at Step 3(b)iiiA. Then there are still at least  $Y_n - 2k\bar{C} - (j-1)$  hospitals with a vacant position and not listed by any couple member at the end of Round  $j-1$ . This follows since at most  $j-1$  hospitals have had their positions filled at Rounds  $1, \dots, j-1$  among those hospitals that are not listed on any single doctor preference at the end of Algorithm 3. Following the steps analogous to those leading to inequality (18), we can compute that Round  $j$ , initiated by some single doctor in  $B^j$ , ends at Step 3(b)iiiA with probability of at least

$$1 - \frac{2k\bar{C}}{\frac{Y_n - 2k\bar{C} - (j-1)}{r} + 2k\bar{C}} > 1 - \frac{2k\bar{C}}{\frac{E[Y_n]/2 - 2k\bar{C} - (j-1)}{r} + 2k\bar{C}}.$$

There are at most  $2\bar{C}$  rounds in Step 3 because at most  $2\bar{C}$  single doctors can be displaced by

couples in Step 2, so  $|B| \leq 2\bar{C}$ . Hence Algorithm 4 succeeds with conditional probability of at least

$$\begin{aligned} \prod_{j=1}^{2\bar{C}} \left( 1 - \frac{2k\bar{C}}{\frac{E[Y_n]/2 - 2k\bar{C} - (j-1)}{r} + 2k\bar{C}} \right) &\geq \left( 1 - \frac{2k\bar{C}}{\frac{E[Y_n]/2 - 2k\bar{C} - (2\bar{C}-1)}{r} + 2k\bar{C}} \right)^{2\bar{C}} \\ &\geq \left( 1 - \frac{2k\bar{C}}{E[Y_n]/4r} \right)^{2\bar{C}}, \end{aligned} \quad (19)$$

where the first inequality follows from Lemma 2, the assumption that  $n$  is sufficiently large and each  $j \leq 2\bar{C}$ , and the second inequality holds since  $E[Y_n]/2 - 4k\bar{C} \geq E[Y_n]/4 > 0$ , which follows from Lemma 2 and the assumption that  $n$  is sufficiently large.

As a result, relations (17) and (19) imply

$$\begin{aligned} &\Pr \left[ \text{Algorithm 4 succeeds} \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu \right] \\ &\geq \left( 1 - \frac{2k\bar{C}r}{n} \right)^{2k\bar{C}} \cdot \left( 1 - \frac{8rk\bar{C}}{E[Y_n]} \right)^{2\bar{C}}. \end{aligned}$$

□

We utilize the following mathematical result (see Lemma 4.4 of Immorlica and Mahdian (2005) for a proof).

**Lemma 4.**  $\text{Var}[Y_n] \leq E[Y_n]$  for every  $n \in \mathbb{N}$ .

### Step 7: Proof of Theorem 1

*Proof of Theorem 1.* We obtain that

$$\begin{aligned} \Pr \left[ Y_n \leq \frac{E[Y_n]}{2} \right] &\leq \Pr \left[ Y_n \leq \frac{E[Y_n]}{2} \right] + \Pr \left[ Y_n \geq \frac{3E[Y_n]}{2} \right] \\ &= \Pr \left[ |Y_n - E[Y_n]| \geq \frac{E[Y_n]}{2} \right] \leq \frac{\text{Var}[Y_n]}{(E[Y_n]/2)^2} \leq \frac{4}{E[Y_n]}, \end{aligned} \quad (20)$$

where the first inequality holds since any probability is nonnegative, the equality is an identity, the second inequality results from Chebyshev inequality, and the last inequality follows from Lemma 4.

By Lemma 3 we have

$$\begin{aligned} & \Pr \left[ \text{Algorithm 4 succeeds} \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu \right] \\ & \geq \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{8rk\bar{C}}{E[Y_n]}\right)^{2\bar{C}}. \end{aligned}$$

This inequality holds for any matching  $\mu$  that is produced at the end of Algorithm 3. Therefore, we have the same lower bound for the probability conditional on  $Y_n > E[Y_n]/2$  but not on  $\mu$ . That is,

$$\Pr \left[ \text{Algorithm 4 succeeds} \mid Y_n > \frac{E[Y_n]}{2} \right] \geq \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{8rk\bar{C}}{E[Y_n]}\right)^{2\bar{C}}. \quad (21)$$

Thus we obtain

$$\begin{aligned} \Pr [\text{Algorithm 4 succeeds}] & \geq \Pr \left[ Y_n > \frac{E[Y_n]}{2} \right] \cdot \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{8rk\bar{C}}{E[Y_n]}\right)^{2\bar{C}} \\ & \geq \left(1 - \frac{4}{E[Y_n]}\right) \cdot \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{8rk\bar{C}}{E[Y_n]}\right)^{2\bar{C}} \\ & \geq \left(1 - \frac{8e^{16\lambda k}}{n}\right) \cdot \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{16rk\bar{C}e^{16\lambda k}}{n}\right)^{2\bar{C}}, \quad (22) \end{aligned}$$

where the first inequality follows from the fact that probabilities are non-negative and (21), the second inequality results from (20), and the last inequality is obtained by Lemma 2.

The first term on the right-hand side of inequality (22),  $\left(1 - \frac{8e^{16\lambda k}}{n}\right)$ , converges to one as long as  $n \rightarrow \infty$ . Consider the second term,  $\left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}}$ . Recall there exists  $b > 0$  such that  $\bar{C} < bn^a$  for any  $n$ . Thus, for any  $n$ ,

$$\begin{aligned} \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} & > \left(1 - \frac{2kbn^a r}{n}\right)^{2kbn^a} \\ & = \left(1 - \frac{2kbr}{n^{1-a}}\right)^{2kbn^{1-a}n^{2a-1}} \\ & \geq (e^{-2kbr})^{2kbn^{2a-1}}. \quad (23) \end{aligned}$$

where the last line follows from the inequality that  $(1 - \frac{\alpha}{x})^x \geq e^{-\alpha}$  when  $x, \alpha > 0$ . Since  $a \in [0, 1/2)$ , the term  $n^{2a-1} \rightarrow 0$  as  $n \rightarrow \infty$  and hence the last expression of inequality (23) converges to one as

$n \rightarrow \infty$ . Similarly, the third term of the right-hand side of inequality (22),  $\left(1 - \frac{16rk\bar{C}e^{16\lambda k}}{n}\right)^{2\bar{C}}$ , can be bounded as

$$\begin{aligned} \left(1 - \frac{16rk\bar{C}e^{16\lambda k}}{n}\right)^{2\bar{C}} &> \left(1 - \frac{16rkbn^ae^{16\lambda k}}{n}\right)^{2bn^a} \\ &= \left(1 - \frac{16rkbe^{16\lambda k}}{n^{1-a}}\right)^{2bn^{1-a}n^{2a-1}} \\ &\geq (e^{-16rkbe^{16\lambda k}})^{2bn^{2a-1}}. \end{aligned} \quad (24)$$

Because  $a \in [0, 1/2)$ , the last expression of inequality (24) converges to one as  $n \rightarrow \infty$ , which concludes the proof.  $\square$

Suppose that the number of couples is bounded along the sequence (which is equivalent to  $a = 0$  in the definition of a regular market). By inequality (22), the probability that there does not exist a stable matching decreases with rate of convergence of  $O(1/n)$  as  $n \rightarrow \infty$ . It is an open question whether this rate of convergence is tight.

## A.4 Proof of Theorem 2

We consider incentives for single doctors, couples, and hospitals in the following subsections and then combine the results to obtain the proof of Theorem 2. Let  $\Pr[E]$  be the probability of the event  $E$ .

### A.4.1 Definition of the mechanism $\varphi$

**Algorithm 5.** MECHANISM  $\varphi$

(1) Initialization:

Let matching  $\mu$  be the output of the deferred acceptance algorithm in the sub-market without couples. Set  $C^0 = C$ ,  $B^0 = \emptyset$ ,  $i = 0$ .

(2) Iterate through couples:

- (a) If  $C^i$  is empty, then go to Step 3. Otherwise, pick some couple  $c = (f, m) \in C^i$  according to some prespecified deterministic rule. Let  $C^{i+1} = C^i \setminus c$  and  $B^{i+1} = B^i$ , and increment  $i$  by one.
- (b) Let couple  $c$  apply to their most preferred pair of hospitals  $(h, h') \in H \times H$  that has not rejected them yet.

- i. If such a hospital (pair) does not exist, modify matching  $\mu$  such that couple  $c$  is unassigned and then go to Step 2a.
- ii. If such a hospital (pair) exists,
  - A. If  $h = h' \neq \emptyset$  and  $\{f, m\} \subseteq Ch_h(\mu(h) \cup c)$ , then modify matching  $\mu$  by assigning  $(f, m)$  to hospital  $h$  and having  $h$  reject

$$(\mu(h) \cup f \cup m) \setminus Ch_h(\mu(h) \cup c).$$

Add the rejected single doctors (if any) to  $B^i$ . If a member of a couple  $c$  is rejected, then have the other member of  $c$  withdraw from the currently matched hospital (if any), and update  $C^i$  by adding  $c$ . Then go to Step 2a.

- B. If  $h \neq h'$ ,  $f \in Ch_h(\mu(h) \cup f)$ , and  $m \in Ch_{h'}(\mu(h') \cup m)$ , then modify matching  $\mu$  by assigning  $f$  to  $h$  and  $m$  to  $h'$ , having hospital  $h$  reject

$$(\mu(h) \cup f) \setminus Ch_h(\mu(h) \cup f),$$

and having hospital  $h'$  reject

$$(\mu(h') \cup m) \setminus Ch_{h'}(\mu(h') \cup m).$$

Add the rejected single doctors (if any) to  $B^i$ . If a member of a couple  $c$  is rejected, then have the other member of  $c$  withdraw from the currently matched hospital (if any), and update  $C^i$  by adding  $c$ . Go to Step 2a.

- C. Otherwise, let hospital  $h$  and hospital  $h'$  reject the application by couple  $c$  and go to Step 2b.

(3) *Iterate through rejected single doctors:*

- (a) If  $B^i$  and  $C^i$  are both empty, then terminate the algorithm.
- (b) Otherwise, if  $B^i$  is empty and  $C^i$  is not, then go to Step 2.
- (c) Otherwise, pick some single doctor  $s$  in  $B^i$ . Let  $B^{i+1} = B^i \setminus s$  and  $C^{i+1}$ , and increment  $i$  by one.

*Iterate through the rank order lists of single doctors:*

- i. If single doctor  $s$  has applied to every acceptable hospital, then modify matching  $\mu$  such that  $s$  is unassigned and go to Step 3a.

- ii. If not, then let  $\hat{h}$  be the most preferred hospital ranked by single doctor  $s$  among those which  $s$  has not yet applied to while running the doctor-proposing deferred acceptance algorithm or previously within this algorithm, and:
  - A. If hospital  $\hat{h}$  prefers not to keep single doctor  $s$ , that is, if  $s \notin Ch_{\hat{h}}(\mu(\hat{h}) \cup s)$ , then let  $\hat{h}$  reject  $s$  and go to Step 3(c)i.
  - B. If hospital  $\hat{h}$  prefers to keep single doctor  $s$ , that is, if  $s \in Ch_{\hat{h}}(\mu(\hat{h}) \cup s)$ , then modify matching  $\mu$  such that  $s$  is assigned to  $\hat{h}$  and  $\hat{h}$  rejects doctor(s)

$$(\mu(\hat{h}) \cup s) \setminus Ch_{\hat{h}}(\mu(\hat{h}) \cup s).$$

If a single doctor is rejected, denote this rejected doctor  $s$  and go to Step 3(c)i. If a member of a couple  $c$  is rejected, then have the other member of  $c$  withdraw from the currently matched hospital (if any), update  $C^i$  by adding  $c$ , and go to Step 3a.

This algorithm terminates in a finite number of steps. We define  $\varphi$  to be the mechanism that, given all reported preferences and capacities, produces the matching reached at the end of this algorithm.

#### A.4.2 Incentives for Single Doctors

Fix the preference  $R_s$  of single doctor  $s$  that is realized from the distribution described in Section 4.1 (we simply say “ $R_s$  is realized” in the remainder of the paper whenever there is no confusion). Define  $\varphi(R'_s)$  to be a random variable which is the matching produced by mechanism  $\varphi$ , where a single doctor  $s$  reports  $R'_s$  and the reported preferences of all other doctors and hospitals are stochastic following Section 4.1. Our first incentive result for  $\varphi$  is about preference manipulations by single doctors.

**Proposition 1.** *Given  $\delta > 0$ , there exists an  $n_0$  such that for every  $n > n_0$ , in random market  $\tilde{\Gamma}^n$ , for any single doctor  $s \in S^n$ , and every pair  $R_s$  and  $R'_s$  of preferences,*

$$\Pr [\varphi(R'_s)(s) P_s \varphi(R_s)(s)] < \delta.$$

*Proof.* Given random market  $\tilde{\Gamma} = (H, S, C, \tilde{\Sigma}_H, k, \mathcal{P}, \mathcal{Q}, \rho)$  that follows assumptions in Section 4, define  $\psi(R'_s)$  to be a random variable which is the matching produced by the doctor-proposing deferred acceptance algorithm on the sub-market  $(H, S, \tilde{\Sigma}_H, k, \mathcal{P})$  that consists of only hospitals and single doctors, where single doctor  $s$  reports  $R'_s$ .

Let us begin by recalling a classic result that the doctor-proposing deferred acceptance algorithm is strategy-proof for single doctors:

**Result 1.** (*Dubins and Freedman, 1981; Roth, 1982*)

$$\psi(R_s)(s)R_s\psi(R'_s)(s),$$

for any single doctor  $s \in S$  and for any preference relations  $R_s$  and  $R'_s$  (and for all realized preferences of other agents).

**Lemma 5.** For any sufficiently large  $n$ , in random market  $\tilde{\Gamma}^n$ , for any single doctor  $s \in S^n$ , preference  $R'_s$ , and matching  $\mu$ ,

$$\begin{aligned} & \Pr \left[ \psi(R'_s)(s) = \varphi(R'_s)(s) \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu \right] \\ & \geq \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]}\right)^{2\bar{C}}, \end{aligned}$$

if the conditioning event has a strictly positive probability.

*Proof.* First, consider the event that all couples are tentatively matched without Algorithm 4 being terminated at Step 2(b)ii so that the algorithm reaches Step 3, and no couple applies to hospital  $\psi(R'_s)(s)$  during Step 2. For that event to happen it is enough for the following event to happen: for any two doctors  $d, d' \in F \cup M$  with  $d \neq d'$ , there is no hospital  $h \in H$  that is listed by both  $d$  and  $d'$  as an acceptable hospital and no  $d \in F \cup M$  draws  $\psi(R'_s)(s)$  from  $\mathcal{Q}^n$ . Following the steps leading to expression (17), this happens with at least probability

$$\left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}}. \tag{25}$$

Next, consider what happens when we assign single doctors in the set  $B$ , conditional on  $Y_n > \frac{E[Y_n]}{2}$ , Algorithm 3 produces matching  $\mu$ , and that all couples are tentatively matched without the algorithm terminating at Step 2(b)ii.

Observe that  $p_{h'} \geq p_h/r$  for any  $h, h' \in H^n$  by assumption. Also observe that there are at most  $2\bar{C}$  hospitals that are tentatively matched to a couple member  $d \in F \cup M$  while there are at least  $Y_n - 2\bar{C} - 1$  hospitals with vacant positions at the beginning of Step 3 (since there are at least  $Y_n - 1$  hospitals with vacant positions at the beginning of Step 2 (the  $-1$  accounts for a position possibly filled by  $s$ ) and at most  $2\bar{C}$  hospitals have their positions filled in Step 2). Following the steps analogous to those leading to inequality (18), we can compute that Round 1 of Step 3 ends at 3(b)iiiA as a single doctor applies to some hospital with vacant positions (rather than applying to a hospital that

is tentatively matched to a couple member), with probability at least

$$1 - \frac{2\bar{C} + 1}{\frac{Y_n - 2\bar{C} - 1}{r} + 2\bar{C} + 1} > 1 - \frac{2\bar{C} + 1}{\frac{E[Y_n]/2 - 2\bar{C} - 1}{r} + 2\bar{C} + 1}.$$

Now assume that all Rounds  $1, \dots, j - 1$  end at Step 3(b)iiiA. Then there are still at least  $Y_n - 2\bar{C} - (j - 1) - 1$  hospitals with a vacant position. This follows since at most  $j - 1$  hospitals have had their positions filled at Rounds  $1, \dots, j - 1$  among those hospitals that are not listed on any single doctor preference at the end of Algorithm 3. Following the steps analogous to those leading to inequality (18), we can compute that Round  $j$  initiated by some single doctor in  $B^j$  ends at Step 3(b)iiiA with probability of at least

$$1 - \frac{2\bar{C} + 1}{\frac{E[Y_n]/2 - 2\bar{C} - (j-1) - 1}{r} + 2\bar{C} + 1}.$$

Since there are at most  $2\bar{C}$  rounds (because this is the maximum number of single doctors who could have been displaced by couples in Step 2), Algorithm 4 succeeds with conditional probability of at least

$$\left(1 - \frac{2\bar{C} + 1}{E[Y_n]/4r}\right)^{2\bar{C}}. \quad (26)$$

Recall that the algorithm underlying  $\varphi$  coincides with the sequential couples algorithm if the latter succeeds. This fact together with the arguments leading to expressions (25) and (26) imply

$$\begin{aligned} & \Pr \left[ \psi(R'_s)(s) = \varphi(R'_s)(s) \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu \right] \\ & \geq \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]}\right)^{2\bar{C}}. \end{aligned}$$

□

To prove Proposition 1 first note that, by Result 1 we have

$$\Pr[\varphi(R_s)(s)R_s\varphi(R'_s)(s)] \geq \Pr[\psi(R_s)(s) = \varphi(R_s)(s) \text{ and } \psi(R'_s)(s) = \varphi(R'_s)(s)]. \quad (27)$$

By Lemma 5 we have

$$\begin{aligned} & \Pr \left[ \psi(R'_s)(s) = \varphi(R'_s)(s) \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu \right] \\ & \geq \left( 1 - \frac{2k\bar{C}r}{n} \right)^{2k\bar{C}} \cdot \left( 1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]} \right)^{2\bar{C}}. \end{aligned}$$

This inequality holds for any matching  $\mu$  that is produced at the end of Algorithm 3. Therefore, we have the same lower bound for the probability conditional on  $Y_n > E[Y_n]/2$  but not on  $\mu$ . That is,

$$\begin{aligned} & \Pr \left[ \psi(R'_s)(s) = \varphi(R'_s)(s) \mid Y_n > \frac{E[Y_n]}{2} \right] \\ & \geq \left( 1 - \frac{2k\bar{C}r}{n} \right)^{2k\bar{C}} \cdot \left( 1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]} \right)^{2\bar{C}}. \end{aligned} \tag{28}$$

By the fact that probabilities are nonnegative and inequalities (28) and then (20), we obtain

$$\begin{aligned} & \Pr [\psi(R'_s)(s) = \varphi(R'_s)(s)] \\ & \geq \Pr \left[ Y_n > \frac{E[Y_n]}{2} \right] \cdot \left( 1 - \frac{2k\bar{C}r}{n} \right)^{2k\bar{C}} \cdot \left( 1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]} \right)^{2\bar{C}} \\ & \geq \left( 1 - \frac{4}{E[Y_n]} \right) \cdot \left( 1 - \frac{2k\bar{C}r}{n} \right)^{2k\bar{C}} \cdot \left( 1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]} \right)^{2\bar{C}}. \end{aligned}$$

By Lemma 2, the last expression converges to one as  $n \rightarrow \infty$ . By an analogous argument, we can establish similar convergence for  $R_s$ . Therefore the right hand side of inequality (27) converges to one.<sup>32</sup> This fact completes the proof of Proposition 1. □

### A.4.3 Incentives for Couples

Our next incentive result is about preference manipulations by couples. Fix the preferences  $R_f$  and  $R_m$  of couple  $c = (f, m)$  that are realized from the distribution described in Section 4.1 (we simply say “ $(R_f, R_m)$  is realized” in the remainder of the paper whenever there is no confusion). Define  $\varphi(R'_c)$  to be a random variable which is the matching produced by mechanism  $\varphi$ , where a couple  $c$  reports a preference relation  $R'_c$  and the reported preferences of all other doctors and hospitals are stochastic following Section 4.1.

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<sup>32</sup>Here we use the fact that if sequences of events  $(F_n)_{n=1}^\infty$  and  $(G_n)_{n=1}^\infty$  satisfy  $\Pr(F_n) \rightarrow 1$  and  $\Pr(G_n) \rightarrow 1$  as  $n \rightarrow \infty$ , then  $\Pr(F_n \cap G_n) \rightarrow 1$  as  $n \rightarrow \infty$ .

**Proposition 2.** *Given  $\delta > 0$ , there exists  $n_0$  such that the following holds: For any  $n > n_0$ , any  $c = (f, m) \in C^n$  and any preference relations  $R_c$  and  $R'_c$ ,*

$$\Pr[\varphi(R'_c)(c)P_c\varphi(R_c)(c)] < \delta.$$

*Proof.* Let  $R_c$  and  $R'_c$  be arbitrary preferences of couple  $c$ . Denote by  $\hat{\varphi}(R'_c)$  the first tentative assignments of couple  $c$  in Algorithm 5 under report  $R'_c$ . Note that in that step, for any  $R'_c$ , couples receive their most preferred hospitals according to  $R'_c$  among those that choose them if  $c$  applies. As a result,  $\hat{\varphi}(R_c)(c)R_c\hat{\varphi}(R'_c)(c)$ . Therefore, it suffices to demonstrate that

$$\Pr[\hat{\varphi}(R_c)(c) = \varphi(R_c)(c), \hat{\varphi}(R'_c)(c) = \varphi(R'_c)(c)] > 1 - \delta.$$

We begin by proving a result similar to Lemma 5.

**Lemma 6.** *For any sufficiently large  $n$ ,  $R'_c$  and  $\mu$ ,*

$$\begin{aligned} & \Pr \left[ \hat{\varphi}(R'_c)(c) = \varphi(R'_c)(c) \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu \right] \\ & \geq \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{8r\bar{C}}{E[Y_n]}\right)^{2\bar{C}} \end{aligned} \quad (29)$$

*if the conditioning event has a strictly positive probability.*

*Proof.* First, consider the event that all couples except for  $c$  are tentatively matched without displacing members of any other couple. Following steps analogous to those leading to expression (17), this happens with at least probability

$$\left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}}. \quad (30)$$

Next, we consider what happens when we assign single doctors in the set  $B$ , conditional on  $Y_n > \frac{E[Y_n]}{2}$  and the event that all couples except for  $c$  are tentatively matched without displacing members of any other couple. Following steps analogous to those leading up to inequality (19), the set of single doctors in  $B$  do not displace any assigned couples with conditional probability of at least

$$\left(1 - \frac{2\bar{C}}{E[Y_n]/4r}\right)^{2\bar{C}}. \quad (31)$$

By relations (30) and (31) and rearranging terms we obtain inequality (29), completing the proof.  $\square$

To prove Proposition 2 recall that

$$\Pr[\varphi(R_c)(c)R_c\varphi(R'_c)(c)] \geq \Pr[\hat{\varphi}(R_c)(c) = \varphi(R_c)(c), \hat{\varphi}(R'_c)(c) = \varphi(R'_c)(c)]. \quad (32)$$

By Lemma 6 we have

$$\begin{aligned} \Pr[\hat{\varphi}(R'_c)(c) = \varphi(R'_c)(c) \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu] \\ \geq \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{8r\bar{C}}{E[Y_n]}\right)^{2\bar{C}}. \end{aligned}$$

This inequality holds for any matching  $\mu$  that is produced at the end of Algorithm 3. Therefore, we have the same lower bound for probability conditional on  $Y_n > E[Y_n]/2$  but not on  $\mu$ . That is,

$$\Pr\left[\hat{\varphi}(R'_c)(c) = \varphi(R'_c)(c) \mid Y_n > \frac{E[Y_n]}{2}\right] \geq \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{8r\bar{C}}{E[Y_n]}\right)^{2\bar{C}}. \quad (33)$$

By the fact that probabilities are nonnegative and inequalities (33) and then (20), we obtain

$$\begin{aligned} \Pr[\hat{\varphi}(R'_c)(c) = \varphi(R'_c)(c)] &\geq \Pr\left[Y_n > \frac{E[Y_n]}{2}\right] \cdot \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]}\right)^{2\bar{C}} \\ &\geq \left(1 - \frac{4}{E[Y_n]}\right) \cdot \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]}\right)^{2\bar{C}}. \end{aligned}$$

By Lemma 2, the last expression converges to one as  $n \rightarrow \infty$ . By an analogous argument, we can establish similar convergence for  $R_c$ . Therefore the right hand side of inequality (32) converges to one. This fact completes the proof of Proposition 2.  $\square$

#### A.4.4 Incentives for Hospitals

Define  $\varphi(\succeq'_h)$  to be a random variable which is the matching produced by mechanism  $\varphi$  when a hospital reports responsive preference  $\succeq'_h$  and the reported preferences of all other hospitals and doctors are stochastic following Section 4.1.

**Proposition 3.** *Given  $\delta > 0$ , there exists an  $n_0$  such that for any  $n > n_0$ , in random market  $\tilde{\Gamma}^n$ , for all hospitals  $h \in H^n$ , for any pair of preferences  $\succeq_h$  and  $\succeq'_h$ ,*

$$\Pr[\varphi(\succeq'_h)(h) \succ_h \varphi(\succeq_h)(h)] < \delta.$$

*Proof.* Define  $\psi(\succeq'_h)$  to be a random variable which is the matching produced by the doctor-proposing deferred acceptance algorithm where the only doctors in the market are single, and hospital  $h$  reports  $\succeq'_h$  while the preferences of doctors are stochastic (and hence the matching is a random variable).

**Lemma 7.** *Given  $\delta' > 0$ , there exists an  $n_0$  such that for any  $n > n_0$ , for all  $h \in H^n$ ,  $\succeq_h$  and  $\succeq'_h$ ,*

$$\Pr[\psi(\succeq_h)(h) \succeq_h \psi(\succeq'_h)(h)] > 1 - \delta'.$$

*Proof.* Lemma 10 of Kojima and Pathak (2009) implies this result.  $\square$

**Lemma 8.** *For any sufficiently large  $n$ ,  $h \in H^n$ ,  $\succeq'_h$  and  $\mu$ ,*

$$\begin{aligned} & \Pr \left[ \psi(\succeq'_h)(h) = \varphi(\succeq'_h)(h) \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu \right] \\ & \geq \left( 1 - \frac{2k\bar{C}r}{n} \right)^{2k\bar{C}} \cdot \left( 1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]} \right)^{2\bar{C}} \end{aligned}$$

*if the conditioning event has a strictly positive probability.*

*Proof.* The proof is analogous to the one for Lemma 5 where we replace  $\varphi(R'_s)(s)$  with hospital  $h$  throughout, so we omit it.  $\square$

To complete the proof, observe that

$$\Pr[\varphi(\succeq_h)(h) \succeq_h \varphi(\succeq'_h)(h)] \geq \Pr[\varphi(\succeq_h)(h) = \psi(\succeq_h)(h), \psi(\succeq_h)(h) \succeq_h \psi(\succeq'_h)(h), \psi(\succeq'_h)(h) = \varphi(\succeq'_h)(h)].$$

By Lemma 8 we have

$$\begin{aligned} & \Pr \left[ \psi(\succeq'_h)(h) = \varphi(\succeq'_h)(h) \mid Y_n > \frac{E[Y_n]}{2} \text{ and Algorithm 3 produces } \mu \right] \\ & \geq \left( 1 - \frac{2k\bar{C}r}{n} \right)^{2k\bar{C}} \cdot \left( 1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]} \right)^{2\bar{C}}. \end{aligned}$$

This inequality holds for any matching  $\mu$  that is produced at the end of Algorithm 3. Therefore, we have the same lower bound for the probability conditional on  $Y_n > E[Y_n]/2$  but not on matching  $\mu$ . That is,

$$\Pr \left[ \psi(\succeq'_h)(h) = \varphi(\succeq'_h)(h) \mid Y_n > \frac{E[Y_n]}{2} \right] \geq \left( 1 - \frac{2k\bar{C}r}{n} \right)^{2k\bar{C}} \cdot \left( 1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]} \right)^{2\bar{C}}. \quad (34)$$

By the fact that probabilities are nonnegative and inequalities (34) and then (20), we obtain

$$\begin{aligned} \Pr[\psi(\succeq'_h)(h) = \varphi(\succeq'_h)(h)] &\geq \Pr\left[Y_n > \frac{E[Y_n]}{2}\right] \cdot \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]}\right)^{2\bar{C}} \\ &\geq \left(1 - \frac{4}{E[Y_n]}\right) \cdot \left(1 - \frac{2k\bar{C}r}{n}\right)^{2k\bar{C}} \cdot \left(1 - \frac{4r(2\bar{C} + 1)}{E[Y_n]}\right)^{2\bar{C}}. \end{aligned}$$

By Lemma 2, the last expression converges to one as  $n \rightarrow \infty$ . A similar convergence result can be obtained for  $\succeq_h$  in lieu of  $\succeq'_h$ . These inequalities and Lemma 7 complete the proof of Proposition 3.

#### A.4.5 Proof of Theorem 2

Fix  $\varepsilon > 0$ . By Propositions 1, 2 and 3, there exists  $n_0$  such that for any random market  $\tilde{\Gamma}^n$  in the sequence with  $n > n_0$ , any player, any of her true and misreported preferences, and any other report of preferences, the probability that she is made better off by that misreporting of preferences when every other player is reporting true preferences is less than

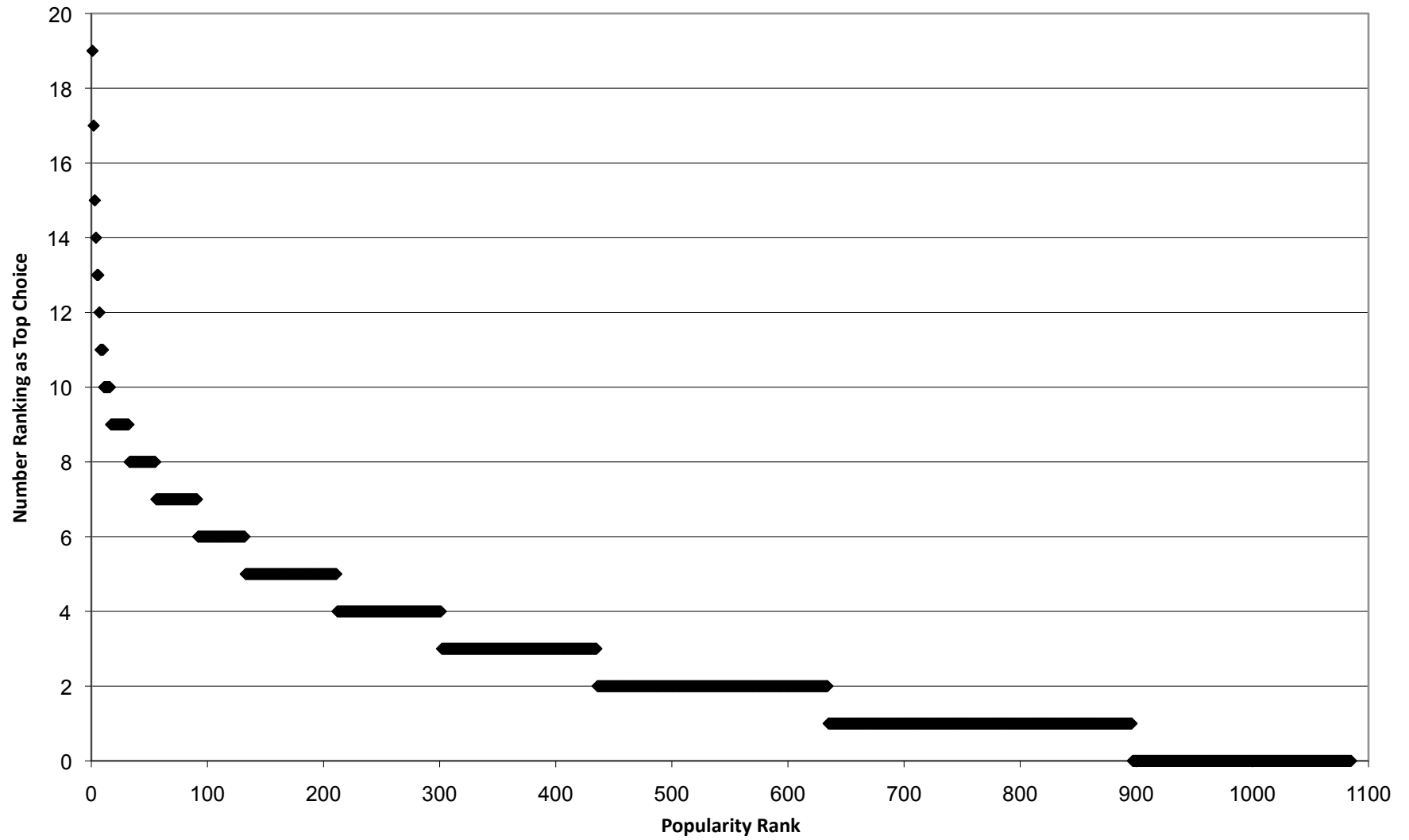
$$\delta = \varepsilon \left( \max\left\{ \sup_{n \in \mathbb{N}, D' \subseteq D^n, h \in H^n, u_h \in \mathcal{U}_h} u_h(D'), \sup_{n \in \mathbb{N}, s \in S^n, h \in H^n, u_s \in \mathcal{U}_s} u_s(h), \sup_{n \in \mathbb{N}, c \in C^n, h, h' \in H^n, u_c \in \mathcal{U}_c} u_c(h, h') \right\} \right)^{-1},$$

where this bound is well-defined and positive by Condition (5) of Definition 2. This observation immediately shows that

$$E[u_i(\varphi_i(\sigma_i(u_i), \sigma_{-i}^*(u_{-i})))] \leq E[u_i(\varphi_i(\sigma^*(u)))] + \varepsilon,$$

for any  $i$ , any type  $u_i$ , and any strategy  $\sigma_i$ , completing the proof.  $\square$

Figure 1: Popularity of Programs as Top Choice (2003)



**Table 1. Summary Statistics for Market for Clinical Psychologists**

	Total	Mean	Min	25th	Median	75th	Max
<i>A. Length of Rank-Order List (ROL)</i>							
Single Applicants	3,010	7.6	1.0	4.0	7.1	10.4	73.1
Couples	19	81.2	7.3	29.4	52.3	115.0	249.9
Distinct Programs Ranked		10.2	2.0	6.4	9.9	13.0	20.9
Programs	1,094	16.7	1.0	7.6	14.3	23.9	80.9
<i>B. Program Capacities</i>							
Capacity	2,721	2.5	1.0	1.0	2.0	3.0	21.4
<i>C. Geographic Similarity of Preferences</i>							
Single Applicants							
# Regions Ranked		2.5	1.0	1.0	2.0	3.3	9.4
Couples							
# Regions Ranked		4.0	1.1	2.6	4.0	4.9	6.9
Fraction of ROL where both Members Rank Same Region		72.7%	29.2%	46.6%	77.3%	97.9%	100.0%

Notes: This table reports descriptive information from the Association of Psychology Postdoctoral and Internship Centers match, averaged over 1999-2007. Single applicants' rank order lists consist of a ranking over hospitals, while couples indicate rankings over program pairs. Distinct programs ranked are the set of distinct programs ranked by each couple member. Programs include only those which have positive capacity. There are 11 regions, corresponding to the first digit of US Zipcodes and Canada.

**Table 2. Comparison of Stable Matchings in Markets with and without Couples**

Matching Market	Applicants						
	Applicant type	Choice Received					
		1st	2nd	3rd	4th	5th+	Unassigned
without couples	single	36.8%	16.9%	10.1%	6.1%	11.2%	18.9%
with couples	single	36.0%	16.6%	10.1%	6.2%	11.6%	19.5%
	couple	18.0%	10.6%	8.7%	5.1%	52.5%	5.2%

Notes: This table reports the choice received in the applicant-optimal stable matching in a market with single applicants and without couples versus a stable matching in the market with couples in the Association of Psychology Postdoctoral and Internship Centers match, averaged over years 1999-2007. An applicant is counted as unassigned even if being unassigned is among her top five choices.

**Table 3. Difference between Stable Matchings in Markets with and without Couples**

Year	Single Applicants Receiving Less Preferred Assignment in Market with Couples		Programs Receiving More Preferred Assignment in Market with Couples	
	Number	Percent	Number	Percent
1999	35	1.2%	45	4.2%
2000	78	2.7%	92	8.4%
2001	86	3.0%	95	8.6%
2002	54	1.9%	62	5.8%
2003	51	1.7%	62	5.7%
2004	65	2.2%	73	6.8%
2005	53	1.7%	63	5.7%
2006	80	2.5%	78	7.1%
2007	69	2.0%	71	6.3%

Notes: This table reports differences between the applicant-optimal in the market without couples and a stable matching in the market with couples in the Association of Psychology Postdoctoral and Internship Centers match. A program receives a more preferred assignment if there is any responsive representation of its preferences for which the assignment is more preferred. There are no single applicants who receive a more preferred assignment in the market with couples and there are no programs that receive a less preferred assignment (for any responsive representation) in the market with couples.

Figure A1: Number of Programs in Each Geographic Region (2003)



**Table A1. Properties of the Stable Set in the Market without Couples**

Year	Single Applicants Receiving More Preferred Assignment in Applicant-Optimal Stable Matching		Programs Receiving Less Preferred Assignment in Applicant-Optimal Stable Matching	
	Number	Percent	Number	Percent
1999	2	0.1%	2	0.2%
2000	7	0.2%	7	0.6%
2001	8	0.3%	8	0.7%
2002	2	0.1%	2	0.2%
2003	6	0.2%	6	0.6%
2004	7	0.2%	7	0.6%
2005	0	0.0%	0	0.0%
2006	6	0.2%	6	0.5%
2007	10	0.3%	10	0.9%

Notes: This table reports differences between the applicant-optimal and program-optimal stable matching in the Association of Psychology Postdoctoral and Internship Centers matching market without couples. A program receives a less preferred assignment if there is any responsive representation of its preferences for which the assignment is less preferred.