# A Simple Augmentation Method for Matchings with Applications to Streaming Algorithms

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# 7 — Abstract

<sup>8</sup> Given a graph G, it is well known that any maximal matching M in G is at least half the size of <sup>9</sup> a maximum matching  $M^*$ . In this paper, we show that if G is bipartite, then running the Greedy <sup>10</sup> matching algorithm on a sampled subgraph of G produces enough additional edges that can be <sup>11</sup> used to augment M such that the resulting matching is of size at least  $(2-\sqrt{2})|M^*| \approx 0.5857|M^*|$ <sup>12</sup> (ignoring lower order terms) with high probability.

The main applications of our method lie in the area of data streaming algorithms, where an 13 algorithm performs few passes over the edges of an *n*-vertex graph while maintaining a memory 14 of size  $O(n \operatorname{polylog} n)$ . Our method immediately yields a very simple two-pass algorithm for 15 MAXIMUM BIPARTITE MATCHING (MBM) with approximation factor 0.5857, which only runs 16 the Greedy matching algorithm in each pass. This slightly improves on the much more involved 17 0.583-approximation algorithm of Esfandiari et al. [ICDMW 2016]. To obtain our main result, we 18 combine our method with a residual sparsity property of the random order Greedy algorithm and 19 give a one-pass random order streaming algorithm for MBM with approximation factor 0.5395. 20 This substantially improves upon the one-pass random order 0.505-approximation algorithm of 21

<sup>22</sup> Konrad et al. [APPROX 2012].

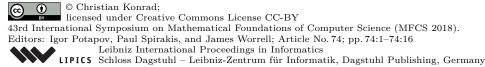
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# <sup>27</sup> **1** Introduction

**Computing Large Matchings.** Given a bipartite graph G = (A, B, E), a matching  $M \subseteq E$ 28 in G is a subset of non-adjacent edges. In this paper, we address the MAXIMUM BIPARTITE 29 MATCHING (MBM) problem, which consists of finding a matching of maximum size. Many 30 classic algorithms for MBM, such as the Hopcroft-Karp algorithm [20] or Edmonds' algorithm 31 [11], as well as many more recent algorithms, first compute an arbitrary matching and then 32 iteratively improve it by finding augmenting paths until it is of maximum size. A good 33 starting point is a *maximal matching*, i.e., a matching that cannot be enlarged by adding an 34 edge outside the matching to it, which is known to be of size at least 1/2 times the size of a 35 maximum matching, i.e., one of maximum size. A maximal matching is for example produced 36 by the GREEDY matching algorithm, which processes the edges of a graph in arbitrary order 37 and adds the current edge to an initially empty matching if the resulting set is still a matching. 38 For an integer  $k \geq 1$ , a (2k+1)-augmenting path  $P = e_1, e_2, e_3, \ldots, e_{2k+1}$  with respect to 39 a matching M is a path of odd length that alternates between edges outside M and edges 40 contained in M such that both  $e_1$  and  $e_{2k+1}$  are incident to vertices that are not matched in 41 M. Since P contains k + 1 edges outside M and k edges of M, removing the matched edges 42



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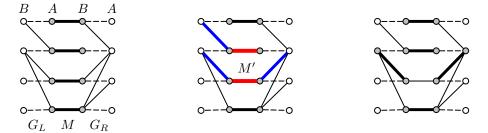
<sup>43</sup> in P from M and inserting the unmatched edges in P into M thus increases the size of M<sup>44</sup> by 1. It is known that a non-maximum matching always admits an augmenting path, and, <sup>45</sup> thus, repeatedly finding one and augmenting eventually yields a maximum matching.

To decrease the number of improvement steps required, one common approach is to compute a large *set of disjoint augmenting paths* and augment along each of them simultaneously. This approach is particularly beneficial when designing algorithms in restricted computational models such as the data streaming model (see below) or various distributed computational models, since typically the number of passes (streaming) or rounds (distributed algorithms) grows linearly with the number of augmentation rounds.

**Our Results.** In this paper, we give a new method that allows us to find a large fraction of 52 disjoint 3-augmenting paths such that, when augmenting along those paths, the resulting 53 matching is of size at least  $(2-\sqrt{2})|M^*| - o(|M^*|) \approx 0.5857|M^*| - o(|M^*|)$  with high 54 probability, where  $M^*$  is a maximum matching (**Theorem 8**). The strength of our method 55 lies both in its simplicity and effectiveness: It only requires running the GREEDY matching 56 algorithm on a random subgraph to produce the necessary edges. Despite its simplicity, it 57 outperforms other more complicated methods and yields improvements over the state-of-the-58 art one- and two-pass data streaming algorithms for matchings (see below). Our method can 59 also be applied repeatedly and for example yields a 3-pass streaming algorithm that also 60 outperforms the currently best 3-pass streaming algorithm known. 61

Applications to Data Streaming Algorithms. While our method can be applied in 62 essentially all computational models that allow an implementation of the GREEDY matching 63 algorithm, it has been designed with the *data streaming model* in mind. Given an *n*-vertex 64 graph G = (V, E), a p-pass, s-space data streaming algorithm processing G performs p passes 65 over the edges E of G (the edges may arrive in arbitrary, potentially adversarial order) 66 while maintaining a memory of size s. Since many graph problems require space  $\Omega(n \log n)$ 67 (observe that storing a large matching already requires this amount of space) [32], research 68 has focussed on the semi-streaming model [16], where a graph streaming algorithm is allowed 69 to use space  $O(n \operatorname{polylog} n)$ . Concerning the MBM problem, Feigenbaum et al. [16] observed 70 that the GREEDY matching algorithm constitutes a one-pass  $\frac{1}{2}$ -approximation semi-streaming 71 algorithm for MBM. Interestingly, despite intense research efforts, no better one-pass 72 streaming algorithms are known, even if space  $O(n^{2-\delta})$  is granted, for any  $\delta > 0$ , while lower 73 bounds only rule out the existence of semi-streaming algorithms with approximation ratio 74 larger than  $1 - 1/e \approx 0.6321$  [22, 18]. Konrad et al. [26] studied minimal extensions to the 75 one-pass semi-streaming model that allow us to improve on GREEDY. They showed that 76 approximation ratios strictly larger than  $\frac{1}{2}$  can be obtained if either the edges of the input 77 graph arrive in uniform random order, or a second pass is granted. More specifically, they 78 gave a symbolic improvement showing that a  $(\frac{1}{2} + 0.005)$ -approximation can be obtained if 79 edges arrive in random order, and a  $(\frac{1}{2} + 0.02)$ -approximation can be achieved if two passes 80 are allowed. Their two-pass result has since been improved by Kale and Tirodkar [21] to 81  $\frac{1}{2} + \frac{1}{16} = \frac{1}{2} + 0.0625$  and independently by Esfandiari et al. to  $\frac{1}{2} + 0.083$  [14]. 82

Our method for finding augmenting paths immediately yields a two-pass semi-streaming 83 algorithm with approximation factor 0.5857 (**Theorem 9**), thus slightly improving over the 84 algorithm of Esfandiari et al. [14]. Our algorithm has constant update time (i.e., the running 85 time between two read operations from the stream) and does not need a post-processing step, 86 while the algorithm of Esfandiari et al. requires the computation of a maximum matching 87 in the post-processing step. Our main result is a one-pass random order semi-streaming 88 algorithm with approximation factor 0.5395 (Theorem 16), showing that more substantial 89 improvements over  $\frac{1}{2}$  than the symbolic improvement given by Konrad et al. [26] are possible 90



**Figure 1** Left: Bipartite graph G = (A, B, E) with maximal matching M. The dotted edges show a perfect matching in G. Matched vertices are grey, free vertices are white. Center: Subset  $M' \subseteq M$  is highlighted in red. The blue edges are produced by the runs of GREEDY on  $G'_L$  and  $G'_R$ . Observe that one 3-augmenting path is found. Right: M after the augmentation.

<sup>91</sup> in the random order scenario. This algorithm is obtained by combining our method for <sup>92</sup> finding augmenting paths with a residual sparsity property of the random order GREEDY <sup>93</sup> matching algorithm (e.g. [25]) that has recently been exploited in various contexts [25, 1, 17].

Techniques. For illustration purposes, consider a bipartite graph G = (A, B, E) that contains a *perfect matching*  $M^*$ , i.e., a matching that matches all vertices, and a maximal matching M with  $|M| = \frac{1}{2}|M^*|$ . It can be seen that  $M \oplus M^* := (M \setminus M^*) \cup (M^* \setminus M)$  forms a set of  $\frac{1}{2}|M^*|$  disjoint 3-augmenting paths. In other words, there exists a matching of size  $\frac{1}{2}|M^*|$  in graph  $G_L := G[A(M) \cup \overline{B(M)}]$ , where A(M) is the set of matched A-vertices, and  $B(M) := B \setminus B(M)$ , and also one of size  $\frac{1}{2}|M^*|$  in  $G_R = G[\overline{A(M)} \cup B(M)]$ , see Figure 1.

We now sample a random subset of edges  $M' \subseteq M$  such that every edge  $e \in M$  is included 100 in M' with probability p. Using an argument by Konrad et al. [26], it follows that when 101 running the GREEDY matching algorithm on the subgraph  $G'_L := G[A(M') \cup B(M)] \subseteq G_L$ , 102 then in expectation a  $\frac{1}{1+p}$  fraction of the vertices A(M') is matched. Observe that if we 103 chose p = 1, then half of the vertices get matched, which is what we expect from the 104 GREEDY matching algorithm. However, if we chose p substantially smaller than 1, then a 105 large fraction of vertices of A(M') is matched. We also apply this argument to subgraph 106  $G'_R := G[B(M') \cup \overline{A(M)}] \subseteq G_R$ , which thus allows us to find large matchings in both 107 subgraphs  $G'_L$  and  $G'_R$  and in turn extract many 3-augmenting paths. Observe that this 108 method directly yields a two-pass semi-streaming algorithm, by computing a maximal 109 matching in the first pass, and augmenting it using the described method in the second pass. 110

The main shortcoming of this method is that the result by Konrad et al. [26] only holds in expectation, which would imply that our result also only holds in expectation. We therefore strengthen their result and prove that a similar version holds with high probability. Our proof models the execution of the algorithm with a Doob martingale and applies Azuma's inequality to obtain a concentration result. We then use our result and additional combinatorial arguments to bound the number of 3-augmenting paths found.

Our one-pass random order streaming algorithm combines our method for finding 3-117 augmenting paths with a *residual sparsity* property of the random order GREEDY algorithm. 118 We run GREEDY on the first  $\frac{1}{\log n}$  fraction of edges in the stream to produce a matching M. 119 The residual sparsity property states that the residual graph  $H = G[V \setminus V(M)]$  contains 120  $O(n \operatorname{polylog} n)$  edges with high probability, which we then collect while processing the 121 remaining edges in the stream. Our main argument is as follows: If |M| is relatively small, 122 then the residual graph H contains a sufficiently large matching. On the other hand, if |M|123 is relatively large (close to a  $\frac{1}{2}$ -approximation), then we can use the remainder of the stream 124 to find 3-augmenting paths using the method described above. 125

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Comparison to Esfandiari et al. [14] and Kale and Tirodkar [21]. The two-pass 126 streaming algorithms of Esfandiari et al. and Kale and Tirodkar proceed similarly in that they 127 compute a maximal matching M in the first pass and then find additional edges in the second 128 pass that are used to augment M. Their algorithms are in fact almost identical and only differ 129 in the post-processing stage. With  $G_L = G[A(M) \cup B(M)]$  and  $G_R = G[B(M) \cup A(M)]$ 130 being as above, they compute incomplete semi-matchings  $S_L$  in  $G_L$  and  $S_R$  in  $G_R$ , i.e., 131 subsets of edges such that every vertex in A(M) (B(M)) is matched at most once in  $S_L$ 132 (resp.  $S_R$ ) and every vertex  $\overline{B(M)}$  (resp.  $\overline{A(M)}$ ) is matched at most k times, for some 133 integer k. Using a Greedy algorithm for computing  $S_L$  and  $S_R$ , it can be seen that a large 134 fraction of vertices A(M) (resp. B(M)) are matched in  $S_L$  (resp.  $S_R$ ). This allows the 135 extraction of multiple 3-augmenting paths. In Kale and Tirodkar, the extraction step is done 136 greedily, which is efficient but leads to a worse approximation factor than in Esfandiari et al. 137 Esfandiari et al. solve an optimization problem in a post-processing phase that allows the 138 extraction of more 3-augmenting paths, which in turn leads to an improved approximation 139 guarantee. Our method is much simpler in this regard, since our additional edges form 140 matchings and it is thus straightforward to extract 3-augmenting paths. 141

Comparison to Konrad et al. [26]. The one-pass random order algorithm by Konrad 142 et al. proceeds as follows: First, run GREEDY on roughly the first third of the edges in the 143 input stream and obtain a matching M. Konrad et al. prove that if GREEDY on the entire 144 input stream produces a matching that is close to a  $\frac{1}{2}$ -approximation, then the matching is 145 built early on, i.e., |M| is relatively large. They then use the remaining part of the stream 146 for finding 3-augmenting paths. To this end, they compute a matching in  $G_L$  on roughly 147 the next third of the edges, and then use the last third to compute a matching in  $G_R$  to 148 complete the 3-augmenting paths. Their method only yields a marginal improvement over 149 1/2 and their result only holds in expectation. 150

Observe that we also argue that the matching M is large, which we achieve by exploiting the residual sparsity property of GREEDY. While Konrad et al. have already processed a third of the edges at this stage, we have only processed a  $\frac{1}{\log(n)}$  fraction, and there are thus more remaining edges to our disposal for finding 3-augmenting paths. Further, our method produces more 3-augmenting paths than the method proposed by Konrad et al.

Further Related Work. Matching problems are the most studied graph problems in the data streaming model. Besides the already mentioned works, algorithms have been designed for weighted matchings (e.g. [16, 29, 33, 9, 31]), multiple passes (e.g. [29, 12, 2]), insertion/deletion streams (e.g. [10, 6, 24, 7, 4, 30]), sparse graphs (e.g. [13, 8]), and other variants of the matching problem [27]. Regarding random order streams, Kapralov et al. [23] showed that the size of a maximum matching can be estimated within a poly-log factor using poly-log space, and a  $(2/3 - \epsilon)$ -approximation can be computed using  $\tilde{O}(n^{3/2})$  space [3].

Outline. We proceed as follows. We first give notation and definitions in Section 2. We then present our method for finding a large set of disjoint 3-augmenting paths in Section 3. Implementation details when implementing our method in the adversarial order streaming model are then discussed in Section 4. In Section 5, we give our one-pass random order algorithm. Finally, we conclude in Section 6 with open problems.

# <sup>168</sup> **2** Preliminaries

<sup>169</sup> Notation. Let G = (A, B, E) be a bipartite graph. We generally use n to denote the number <sup>170</sup> of vertices, i.e., n = |A| + |B|, and m = |E| to denote the number of edges. For a subset of

vertices  $U \subseteq A \cup B$  and a subset of edges  $F \subseteq E$ , we denote the vertex induced subgraph 171 of G by vertices U by G[U], and the edge induced subgraph of G by edges F by G[F]. Let 172 M be a matching in G. We denote by A(M)(B(M)) the vertices of A (resp. B) that are 173 matched by M, and we write  $V(M) = A(M) \cup B(M)$ . Similarly, for an edge  $e \in E$ , we write 174 A(e) to denote its incident A-vertex, B(e) to denote its B vertex, and  $V(e) = \{A(e), B(e)\}$ . 175 The complement of a subset  $A' \subseteq A$   $(B' \subseteq B)$  is denoted by  $\overline{A'} = A \setminus A'$  (resp.  $\overline{B'} = B \setminus B'$ ). 176 The matching number of a graph G, i.e., the size of a maximum matching in G, is denoted 177  $\mu(G)$ . We write opt(G) to denote an arbitrary but fixed maximum matching in G. For two 178 sets X, Y, we write  $X \oplus Y := (X \setminus Y) \cup (Y \setminus X)$  to denote their symmetric difference. For a 179 graph  $G, \Delta(G)$  denotes the maximum degree. 180

Concentration Bounds. In this paper, we will use two concentration bounds. The first one is Azuma's inequality for martingales (we refer the reader to [28] for the an introduction to martingales and Azuma's inequality), and the second is a Chernoff-type bound for weakly dependent random variables.

▶ Theorem 1 (Azuma's Inequality ([5, 28])). Suppose that  $X_0, X_1, X_2, ...$  is a martingale and let  $|X_i - X_{i-1}| \le c_i$  for suitable constants  $c_i$ . Then:

$$\mathbb{P}\left[|X_n - X_0| \ge t\right] \le 2 \exp\left(\frac{-t^2}{2\sum_{i=1}^n c_i^2}\right) \ .$$

▶ **Theorem 2** (Chernoff Bound for Weakly Dependent Variables, e.g. [15]). Let  $X_1, X_2, ..., X_n$  be 0/1 random variables for which there is a  $p \in [0, 1]$  such that for all  $k \in [n]$  the inequality

$$\mathbb{P}\left[X_k = 1 \mid X_1, X_2, \dots, X_{k-1}\right] \le p$$

holds (i.e., the probability of  $X_k = 1$  conditioned on any possible outcome of  $X_1, \ldots, X_{k-1}$  is at most p). Let further  $\mu \ge p \cdot n$ . Then, for every  $\delta > 0$ :

$$\mathbb{P}\left[\sum_{i=1}^{n} X_i \ge (1+\delta)\mu\right] \le \left(\frac{e^{\delta}}{(1+\delta)^{1+\delta}}\right)^{\mu} .$$

We will say that an event occurs with *high probability in variable x*, if the the probability of the event occurring is at least  $1 - x^{-C}$ , for some  $C \ge 1$ . If we do not mention x explicitly, then the high probability statement is in n, the number of vertices of the input graph.

We say that an algorithm is a C-approximation algorithm for MBM if it computes a matching M of size at least  $C \cdot \mu(G) - o(\mu(G))$ .

## **3** Finding a Large Set of Disjoint 3-augmenting Paths

<sup>191</sup> We now present an algorithm that, given a maximal matching M in a bipartite graph <sup>192</sup> G = (A, B, E), finds a set of disjoint 3-augmenting paths  $\mathcal{P}$  by running the GREEDY <sup>193</sup> matching algorithm on a random subgraph of G. The set  $\mathcal{P}$  is such that, when augmenting M<sup>194</sup> along the paths  $\mathcal{P}$ , a matching of size at least  $(2 - \sqrt{2})\mu(G) - o(\mu(G)) \approx 0.5857\mu(G) - o(\mu(G))$ <sup>195</sup> is obtained.

Our algorithm is illustrated in Algorithm 1. For the sake of a clear presentation, the algorithm employs two invocations of GREEDY on two disjoint subgraphs. This is equivalent to invoking GREEDY only once on their union. Our algorithm is parametrized by a sampling probability *p*. To obtain the claimed bound stated above, we will later optimize *p*.

To obtain a better understanding of our algorithm, we first discuss structural properties that help us locate 3-augmenting paths in G with respect to the matching M.

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**Input:** Bipartite graph G = (A, B, E), maximal matching M, parameter 0

- 1. Sample each edge  $e \in M$  with probability p; let M' be the resulting sample
- 2.  $M_L \leftarrow \text{GREEDY}(G[A(M') \cup \overline{B(M)}]); M_R \leftarrow \text{GREEDY}(G[\overline{A(M)} \cup B(M')])$
- **3.**  $\mathcal{P} \leftarrow \{ \text{paths } b'a, ab, ba' \mid b'a \in M_L, ab \in M, ba' \in M_R \}$
- 4. return  $\mathcal{P}$

Algorithm 1. Finding a large set of 3-augmenting paths

Let  $M^*$  be a maximum matching in G and let  $\epsilon$  be such that  $|M| = (\frac{1}{2} + \epsilon)|M^*|$ . Observe first that  $M \oplus M^*$  contains a collection of  $(\frac{1}{2} - \epsilon)|M^*|$  disjoint augmenting paths. Further, observe that the endpoints of each augmenting path are a free vertex in A (i.e., a vertex in  $\overline{A(M)}$ ) and a free vertex in B. Hence, the subgraphs  $G_L := G[A(M) \cup \overline{B(M)}]$  and  $G_R := G[\overline{A(M)} \cup B(M)]$  each contain a matching of size  $(\frac{1}{2} - \epsilon)|M^*|$ . We summarize this in Observation 3:

▶ **Observation 3.** Let  $\epsilon$  be such that  $|M| = (\frac{1}{2} + \epsilon)\mu(G)$ . Then:

$$\min\{\mu(G_L), \mu(G_R)\} \ge (\frac{1}{2} - \epsilon)\mu(G)$$

Suppose now that  $\epsilon$  is small. Further, suppose that we could compute maximum matchings  $M_L^*$  in  $G_L$  and  $M_R^*$  in  $G_R$ . Then for almost every edge  $e \in M$  there are edges  $e_l \in M_L$  and  $e_r \in M_R$  such that  $e_l ee_r$  forms a 3-augmenting path. We will call  $e_l$  a left wing for edge eand  $e_r$  a right wing for edge e.

Our augmentation method should of course not be based on computing maximum 212 matchings themselves. We therefore proceed differently. First, observe that if we computed 213 maximal matchings, i.e.,  $\frac{1}{2}$ -approximations, in  $G_L$  and  $G_R$ , then we may not find any 3-214 augmenting path at all, since it may happen that we find left wings for half of the edges of 215 M, and right wings for the other half. Our strategy therefore is as follows: We first sample a 216 subset of edges  $M' \subseteq M$ , where each edge of M is included in M' with probability p, and we 217 attempt to augment only the edges in M' by computing GREEDY matchings in the subgraphs 218  $G'_L := G[A(M') \cup \overline{B(M)}]$  and  $G'_R := G[\overline{A(M)} \cup B(M')]$ . Konrad et al. [26] proved that, in 219 expectation, the resulting matchings are essentially  $\frac{1}{1+p} \geq \frac{1}{2}$ -approximations, albeit for a 220 slightly different notion of approximation, which is nevertheless suitable for our purposes: 221

▶ Theorem 4 (Konrad et al. [26]). Let G = (U, V, E) be a bipartite graph, and let  $U' \subseteq U$  be such that every vertex  $u \in U$  is included in U' with probability p ( $p \in [0, 1]$ ). Then, for any arbitrary but fixed order in which GREEDY processes the edges, the following holds:

$$\mathbb{E}_{U'}|\operatorname{GREEDY}(G[U'\cup V])| \ge \frac{p}{1+p}\mu(G) \ .$$

Hence, if  $\epsilon$  is close to 0, and p is substantially smaller than 1, then it follows from the previous theorem that a large fraction of the vertices A(M') will be matched by GREEDY in  $G'_L$ , and a large fraction of the vertices of B(M') will be matched by GREEDY in  $G'_R$ . This in turn implies that a substantial amount of edges of M' both have left and a right wings and are thus included in 3-augmenting paths.

Before we make this intuition formal, we point out one shortcoming of applying the previous theorem by Konrad et al. directly. They prove that the resulting matching is large only *in expectation*, which in turn would imply that our result only holds in expectation.

We therefore first strengthen their result and prove that a similar version holds with high probability. To this end, we first prove a technical lemma that is employed in the proof of our strengthened theorem.

▶ Lemma 5. Let G = (U, V, E) be a bipartite graph and let  $u \in U, v \in V$  be arbitrary vertices. Let  $U' \subseteq U$  be such that every vertex  $u \in U$  is included in U' with probability p. Then, for any arbitrary but fixed order in which GREEDY processes the edges, the following holds:

 $0 \leq \mathbb{E}_{U'} |\operatorname{GREEDY}(G[U' \cup V])| - \mathbb{E}_{U'} |\operatorname{GREEDY}(G[(U' \cup V) \setminus \{u, v\}])| \leq 2.$ 

233 **Proof.** First, observe that

$$\mathbb{E}_{U'}|\operatorname{GREEDY}(G[U'\cup V])| - \mathbb{E}_{U'}|\operatorname{GREEDY}(G[(U'\cup V)\setminus \{u,v\}])| =$$

$$\mathbb{E}_{U'}\left(|\operatorname{GREEDY}(G[U'\cup V])| - |\operatorname{GREEDY}(G[(U'\cup V)\setminus \{u,v\}])|\right) .$$

We will prove next that  $0 \leq \text{GREEDY}(G[U' \cup V]) - \text{GREEDY}(G[U' \cup V - \{u, v\}]) \leq 2$ holds for any  $U' \subseteq U$ , which then proves the lemma. We will in fact argue the stronger statement that for any graph G = (V, E) and any vertex  $v \in V$ , the inequality  $0 \leq$ GREEDY $(G) - \text{GREEDY}(G \setminus \{v\}) \leq 1$  holds. The result then follows by applying this statement twice.

Consider thus an arbitrary graph G = (V, E) and a vertex  $v \in V$ . First observe that if GREEDY(G) leaves v unmatched, then GREEDY(G) = GREEDY( $G \setminus \{v\}$ ). If GREEDY(G) matches v, then it is not hard to see that GREEDY(G)  $\oplus$  GREEDY( $G \setminus \{v\}$ ) consists of one alternating path whose one endpoint is v. This further implies that GREEDY( $G \setminus \{v\}$ )  $\leq$ GREEDY(G)  $\leq$  GREEDY( $G \setminus \{v\}$ ) + 1, which completes the proof.

We now give our strengthened version of Theorem 4.

▶ **Theorem 6.** Let G = (U, V, E) be a bipartite graph, and let  $U' \subseteq U$  be such that every vertex  $u \in U$  is included in U' with probability p ( $p \in [0, 1]$ ). Then, for any arbitrary but fixed order in which GREEDY processes the edges, the following holds with probability at least  $1 - (\mu(G))^{-12}$ :

$$|\operatorname{GREEDY}(G[U' \cup V])| \ge \frac{p}{1+p}\mu(G) - o(\mu(G)) .$$

<sup>247</sup> **Proof.** Let  $X := |\text{GREEDY}(G[U' \cup V])|$ . By Theorem 4 we have  $\mathbb{E}X \ge \frac{p}{1+p}\mu(G)$ .

For  $1 \le i \le n$ , let  $Z_i$  be the *i*th edge selected by GREEDY, and let  $Z_i = \bot$  if i > X. Let  $Y_i$  be the Doob martingale induced by the first *i* choices of the algorithm, i.e.,

$$Y_i := \mathbb{E}_{Z_{i+1}, Z_{i+2}, \dots, Z_n} (X \mid Z_1, \dots, Z_i)$$

Observe that the expectation in the previous expression is in itself a random variable, since the expectation is only taken over  $Z_{i+1}, \ldots, Z_n$ , while  $Z_1, \ldots, Z_i$  are random variables. It is not hard to check that the sequence  $(Y_i)_i$  always forms a martingale, independently of the underlying sequence  $Z_i$ . Observe next that  $Y_0 = \mathbb{E}X$  and  $Y_n = X$ . We thus need to show that  $|Y_n - Y_0|$  is small with high probability. To this end, we will apply Azuma's inequality, which requires bounding the differences  $|Y_{i+1} - Y_i|$ , for every *i*, first.

First, observe that  $|Y_{i+1} - Y_i| = 0$  for every  $i \ge X$ . Next, we claim that  $|Y_{i+1} - Y_i| \le 1$ , for every i < X. Indeed, observe that  $Y_i$  is the expected size of the computed matching conditioned on the first *i* choices of the algorithm. We can thus rewrite  $Y_i$  as:

$$Y_i = i + \mathbb{E}_{U'} |\text{GREEDY}(H_i)| ,$$

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where  $H_i := G[(U' \cup V) \setminus \bigcup_{j \leq i} V(Z_j)]$  is the residual graph obtained when removing the 256 vertices incident to the first i selected edges. We thus obtain: 257

where we applied Lemma 5. 260

Next, since  $X \leq \mu(G[U' \cup V]) \leq \mu(G)$ , we have  $|Y_{i+1} - Y_i| \leq 1$  for every  $i \leq \mu(G)$ , and 261  $|Y_{i+1} - Y_i| = 0$  for every  $i > \mu(G)$ . Applying Azuma's Inequality (Theorem 1), we obtain: 262

$$\mathbb{P}\left[|Y_n - Y_0| \ge 5\sqrt{\mu(G)\ln(\mu(G))}\right] \le \mu(G)^{-12} .$$

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Equipped with Theorem 6, we now show that our algorithm finds many disjoint 3-264 augmenting paths, provided that M is close to a  $\frac{1}{2}$ -approximation. 265

**Lemma 7.** Consider Algorithm 1 and suppose that  $|M| = (\frac{1}{2} + \epsilon)\mu(G)$ . Then, with probability at least  $1 - \mu(G)^{-10}$ ,

$$|\mathcal{P}| \ge \mu(G)p\left(\frac{1-2\epsilon}{1+p} - \frac{1}{2} - \epsilon\right) - o(\mu(G))$$

**Proof.** First, by an application of a Chernoff bound, we obtain  $|M'| = p|M| \pm O(\sqrt{|M| \ln(|M|)})$ , 266 with probability at least  $1 - |M|^{-C}$ , for an arbitrarily large constant C. Next, by The-267 orem 6 and Observation 3, with probability at least  $1 - 2(\mu(G))^{-12}$ , we have  $|M_L| \ge$ 268  $\frac{p}{1+p}(\frac{1}{2}-\epsilon)\mu(G) - o(\mu(G)) \text{ and } |M_R| \geq \frac{p}{1+p}(\frac{1}{2}-\epsilon)\mu(G) - o(\mu(G)). \text{ Observe that at most } |M'| - |M_L| \text{ edges of } M' \text{ do not have a left wing, and at most } |M'| - |M_R| \text{ edges of } M' \text{ do not } have a left wing, and at most } |M'| - |M_R| \text{ edges of } M' \text{ do not } have a left wing, and at most } |M'| - |M_R| \text{ edges of } M' \text{ do not } have a left wing, and at most } |M'| - |M_R| \text{ edges of } M' \text{ do not } have a left wing, and at most } |M'| - |M_R| \text{ edges of } M' \text{ do not } have a left wing, and at most } |M'| - |M_R| \text{ edges of } M' \text{ do not } have a left wing, and at most } |M'| - |M_R| \text{ edges of } M' \text{ do not } have a left wing, and at most } \|M'\| - \|M_R\| \text{ edges of } M' \text{ do not } have a \|M'\| - \|M_R\| + \|M_R$ 269 270 have a right wing. Hence, at least  $|M'| - (|M'| - |M_L|) - (|M'| - |M_R|) = |M_L| + |M_R| - |M'|$ 271 edges have both left and right wings and therefore form 3 augmenting paths. We thus obtain: 272

$$\begin{aligned} {}_{273} \quad |\mathcal{P}| &\geq |M_L| + |M_R| - |M'| \geq 2 \cdot \frac{p}{1+p} (\frac{1}{2} - \epsilon) \mu(G) - o(\mu(G)) - p|M| - O(\sqrt{|M|\ln(|M|)}) \\ &\geq 2 \cdot \frac{p}{1+p} (\frac{1}{2} - \epsilon) \mu(G) - p(\frac{1}{2} + \epsilon) \mu(G) - o(\mu(G)) \end{aligned}$$

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$$= \mu(G)p\left(\frac{1-2\epsilon}{1+p} - \frac{1}{2} - \epsilon\right) - o(\mu(G)) .$$

By the union bound, the error is bounded by  $|M|^{-C} + 2(\mu(G))^{-12} \leq (\mu(G))^{-10}$ . 276

We are now ready to prove our main theorem: 277

**Theorem 8.** Let M be a maximal matching. Then, setting  $p = \sqrt{2} - 1$  in Algorithm 1 278 guarantees that M augmented by  $\mathcal{P}$  gives a matching of size at least  $(2-\sqrt{2})\mu(G) - o(\mu(G)) \approx$ 279  $(\frac{1}{2}+0.0857)\mu(G)-o(\mu(G))$  with high probability in  $\mu(G)$ . 280

**Proof.** Observe that the final matching is of size  $|M| + |\mathcal{P}|$ . Let  $\epsilon$  be such that |M| = $(\frac{1}{2}+\epsilon)\mu(G)$ . By Lemma 7, we have

$$|M| + |\mathcal{P}| \geq (\frac{1}{2} + \epsilon)\mu(G) + \mu(G)p\left(\frac{1-2\epsilon}{1+p} - \frac{1}{2} - \epsilon\right) - o(\mu(G)) .$$
(1)

It can be seen that for any value of p, the right side of Inequality 1 is minimized for  $\epsilon = 0$ . On the other hand, for any value of  $\epsilon$ , the value  $p(\epsilon) = \sqrt{\frac{1-2\epsilon}{\frac{1}{2}+\epsilon}} - 1$  maximizes Inequality 1. 285

Using 
$$\epsilon = 0$$
 and  $p(0) = \sqrt{2} - 1$  in Inequality 1 gives  $|M| + |\mathcal{P}| \ge (2 - \sqrt{2})\mu(G) - o(\mu(G))$ .

<sup>287</sup> Multiple augmentation rounds with decreasing values of p allow further improvements. For <sup>288</sup> example, a second round with  $p = \sqrt{\frac{2-\sqrt{2}}{\sqrt{2}-1}} - 1 \approx 0.1892$  guarantees that the resulting <sup>289</sup> matching is of size at least  $0.6067\mu(G) - o(\mu(G))$ . As we will discuss in the next section, this <sup>290</sup> can give a 3-pass streaming algorithm for MBM with approximation factor 0.6067, which <sup>291</sup> slightly improves the 3-pass 0.605-approximation algorithm by Esfandiari et al. [14].

# **4** Adversarial Order Streams

Our method for finding augmenting paths given in Section 3 can directly be implemented in the streaming model. In the first pass, we compute a maximal matching M. If the current edge is added to M, then with probability p we add the edge to M' as well. In the second pass, we run GREEDY on the subgraphs  $G'_L$  and  $G'_R$  and as soon as a 3-augmenting path is completed, we augment M. This can be done with constant update times.

Since we would like our streaming algorithm to succeed with high probability in n, the 298 number of vertices, we need to address the fact that our method as stated in Theorem 8 only 299 succeeds with high probability in  $\mu(G)$ , the size of a maximum matching in G. If  $\mu(G)$  is of 300 size at least, say,  $\Omega(n^{\frac{1}{4}})$ , our method can also gives a high probability result with respect 301 to n. To deal with the case  $\mu(G) = o(n^{\frac{1}{4}})$  we run the 1-pass algorithm of Chitnis et al. [7] 302 in parallel to our algorithm, which computes a subset of edges  $E' \subseteq E$  of size  $O(n^{\frac{1}{2}})$  that 303 contains a maximum matching provided that  $\mu(G) = O(n^{\frac{1}{4}})$ . Observe that after the first 304 pass, we know in which of the two cases we are. We then run the Hopcroft-Karp maximum 305 matching algorithm [20] in time  $O(\sqrt{n} \cdot \sqrt{n}) = O(n)$  on the set of collected edges. To obtain 306 a streaming algorithm with constant update time, we amortize the previous computation 307 during the processing of the second pass, which is possible under the natural assumption 308 that  $m = \Omega(n)$ . This gives the following theorem: 309

**Theorem 9.** There is a two-pass streaming algorithm for MBM with approximation factor  $2 - \sqrt{2} \approx \frac{1}{2} + 0.0857$  that succeeds with high probability (in n). Using one additional pass, a 0.6067-approximation algorithm can be obtained.

# **5** 1-pass Random Order Streaming Algorithm

In this section, we assume that  $\mu(G) = \Omega(n^{\frac{1}{4}})$ . To deal with the case  $\mu(G) = o(n^{\frac{1}{4}})$  we run the algorithm of Chitnis et al. [7] as outlined in Section 4 in parallel and compute and output a maximum matching after processing the stream. We also assume that the input graph has at least  $C_1 \cdot n \log^{C_2} n$  edges, for suitably large constants  $C_1, C_2$ . If this is not the case then we could simply store all edges within the semi-streaming space constraint and compute and output a maximum matching.

Our 1-pass random order streaming algorithm combines our method for finding augmenting paths with a *residual sparsity* property of the random order GREEDY matching algorithm:

▶ Theorem 10 (Residual Sparsity of GREEDY). Suppose that GREEDY processes the edges E of a graph G = (V, E) with m = |E| in uniform random order. Let  $M_i$  be the matching produced by GREEDY after having processed the ith edge. Then:

$$\Delta(G[V \setminus V(M_i)]) = O(\frac{m \log n}{i})$$

with probability  $1 - n^{-12}$  (over the uniform random ordering of the edges).

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- This theorem is implied by a similar theorem concerning the random order GREEDY algorithm for independent sets as given in [25]. Observe that the GREEDY algorithm for matchings on a
- randomly ordered sequence of the edges of a graph G can be seen as the GREEDY algorithm
- for independent sets on a randomly ordered sequence of the vertices of the line graph L(G).
- Our one-pass random order algorithm is parametrized by a probability p, and is illustrated in Algorithm 2. In this listing, we write  $\pi = \pi[1], \pi[2], \ldots, \pi[m]$  to be a uniform random
- in Algorithm 2. In this listing, we write  $\pi = \pi[1], \pi[2], \ldots, \pi[m]$  to be a uniform random ordering of the edges E. For a < b we also write  $\pi[a, b]$  to denote edges  $\pi[a], \pi[a+1], \ldots, \pi[b]$ , and  $\pi(a, b]$  to denote edges  $\pi[a+1], \pi[a+2], \ldots, \pi[b]$ .

**Input:** Bipartite graph G = (A, B, E) with m edges, parameter 0

Let  $\pi = \pi[1], \pi[2], \dots \pi[m]$  be the edges of G in uniform random order

- **1.**  $M \leftarrow \text{Greedy}(\pi[1, \frac{m}{\log n}])$
- **2.** Let  $M' \subseteq M$  be such that every edge of M is included in M' with probability p
- 3. while processing  $\pi(\frac{m}{\log n}, m]$  do in parallel:
  - **a.** Compute set  $E_M$  of edges  $ab \in \pi(\frac{m}{\log n}, m]$  with  $a, b \notin V(M)$ ; if  $|E_M| \ge C \cdot n \log^2 n$ , for some appropriate large constant C, then **abort**
  - **b.**  $M_L \leftarrow \text{GREEDY}(G_L^r)$ , where  $G_L^r$  is the subgraph of G induced by all edges  $\pi(\frac{m}{\log n}, m]$  between A(M') and  $\overline{B(M)}$
  - c.  $M_R \leftarrow \text{GREEDY}(G_R^r)$ , where  $G_R^r$  is the subgraph of G induced by all edges  $\pi(\frac{m}{\log n}, m]$  between  $\overline{A(M)}$  and B(M')
- 4.  $\mathcal{P} \leftarrow \{ \text{paths } b'a, ab, ba' \mid b'a \in M_L, ab \in M, ba' \in M_R \}$
- 5. if  $|\mathcal{P}| \geq \mu(G[E_M])$  then return M augmented by  $\mathcal{P}$ 
  - else return  $M \cup opt(G[E_M])$

# Algorithm 2. One-pass random order matching algorithm

We run GREEDY on the first  $\frac{m}{\log n}$  edges to compute a matching M. Theorem 10 implies that the maximum degree in the residual graph  $H := G[V \setminus V(M)]$  is  $O(\log^2 n)$ . This allows us to collect the entire residual graph (i.e., set  $E_M$ ) within the semi-streaming space bound, since it has  $O(n \log^2 n)$  edges with high probability. We abort if  $|E_M|$  becomes too large. In the next stage, we proceed as in our two-pass algorithm: We sample a subset of edges

<sup>336</sup>  $M' \subseteq M$  and we try to find 3-augmenting paths for M' by computing matchings  $M_L$  and <sup>337</sup>  $M_R$  in the subgraphs  $G_L^r$  and  $G_R^r$ . Ideally we would like to search for left and right wings <sup>338</sup> in the subgraphs  $G_L := G[A(M) \cup \overline{B(M)}]$  and  $G_R := G[\overline{A(M)} \cup B(M)]$ . Since however the <sup>339</sup> first  $\frac{1}{\log n}$  fraction of edges in the stream has already been processed, we can only search for <sup>340</sup> augmenting paths in  $G_L^r$  and  $G_R^r$ . Concentration bounds however allow us to prove that not <sup>341</sup> many important edges have arrived among the first  $\frac{1}{\log n}$  fraction of edges (Lemma 14).

Our analysis is build on the following important observation. Suppose first that the matching M is small, i.e.,  $|M| = \alpha |M^*|$ , for a small value of  $\alpha$ . Then we will argue in the next lemma that a maximum matching in the residual graph is large:

▶ Lemma 11. Let  $\alpha$  be such that  $|M| = \alpha |M^*|$ , and let  $H := G[E_M]$  (=  $G[V \setminus V(M)]$ ) be the residual graph. Then:

$$\mu(H) \ge (1 - 2\alpha)|M^*|$$
.

Proof. Let  $M^*$  be a maximum matching in G. Let  $M_1^* \subseteq M^*$  be those edges of  $M^*$  that share at least one endpoint with an edge in M, and let  $M_2^* = M^* \setminus M_1^*$ . Then  $|M_1^*| \leq 2|M|$ , since each edge of M can only be incident to at most two edges of  $M^*$ . Observe further that  $M_2^* \subseteq E_M$ . Hence:  $\mu(H) \geq |M_2^*| = |M^*| - |M_1^*| \geq |M^*| - 2|M| = (1 - 2\alpha)|M^*|$ .

 $_{349}$  By combining M with a maximum matching in H we obtain the following corollary:

**Some Corollary 12.** Algorithm 2 finds a matching of size at least  $(1-\alpha)|M^*|$  with high probability.

The previous corollary shows that either the matching  $M \cup opt(H)$  is large (if  $\alpha$  is small), or the matching M itself is already reasonably large (if  $\alpha$  is large). This is an important property since we next attempt to augment M, which necessitates that M is already close to  $\frac{1}{2}$ -approximation. For this to succeed, we need to show that  $\mu(G_L^r)$  and  $\mu(G_R^r)$  are large. To this end, let  $\delta$  be such that  $|M| + \mu(G[E_M]) = (\frac{1}{2} + \delta)\mu(G)$ . We will first bound  $\mu(G_L)$ and  $\mu(G_R)$  and then prove a similar bound for  $\mu(G_L^r)$  and  $\mu(G_R^r)$ .

▶ Lemma 13. Suppose that  $|M| + \mu(G[E_M]) = (\frac{1}{2} + \delta)\mu(G)$ . Then:

$$\min\{\mu(G_L), \mu(G_R)\} \ge (\frac{1}{2} - \delta)\mu(G)$$

**Proof.** Let  $M^*$  be a maximum matching in G and let  $M_H^*$  be an arbitrary maximum matching in  $H(=G[E_M])$ . First, it is not hard to see that  $M \cup M_H^*$  is a maximal matching. Next, consider the set of edges  $M^* \oplus (M \cup M_H^*)$ . Since  $|M| + |M_H^*| = (\frac{1}{2} + \delta)\mu(G)$ , the set  $M^* \oplus (M \cup opt(H))$  contains  $(\frac{1}{2} - \delta)\mu(G)$  augmenting paths.

Observe that none of these augmenting paths only contain edges of  $M^*$  and  $M_H^*$ , since this would imply that  $M_H^*$  is not maximum in H. Consider now one such augmenting path P and remove all edges of  $M_H^*$  from P. Then P contains at least one augmenting path that only contains edges from M and  $M^*$ . Applying this argument to all augmenting paths, this proves that there are matchings in  $G_L$  and  $G_R$  of sizes  $(\frac{1}{2} - \delta)\mu(G)$ .

▶ Lemma 14. Suppose that  $|M| + \mu(G[E_M]) = (\frac{1}{2} + \delta)\mu(G)$ . Then, with high probability,

$$\min\{\mu(G_L^r), \mu(G_R^r)\} \ge (1 - \frac{4}{\log n}) \cdot (\frac{1}{2} - \delta)\mu(G)$$

Proof. We only give the argument for  $G_L^r$ , the argument for  $G_R^r$  is identical. Let  $M_L^* = opt(G_L)$ . We will show that most edges of  $M_L^*$  are included in  $\pi(\frac{m}{\log n}, m]$  with high probability. By Lemma 13, we have  $|M_L^*| \ge (\frac{1}{2} - \delta)\mu(G)$ . Let  $e_i$  be the *i*-th edge of  $M_L^*$ , let  $t_i$  be its position in the stream, and let  $Y_i$  be the indicator variable of the event " $t_i \le \frac{m}{\log n}$ ". Our aim is to bound the probabilities  $\mathbb{P}[Y_i = 1 | Y_1, \ldots, Y_{i-1}]$  and then apply the Chernoff bound stated in Theorem 2.

In the following, all our arguments are conditioned on the event " $|E(G[V \setminus V(M)])| = O(n \log^2 n)$ " (without explicitly mentioning it), which we denote by  $E_1$ . This implies that the algorithm does not abort in Line 3a. By the residual sparsity property as stated in Theorem 10,  $E_1$  occurs with probability at least  $1 - n^{-12}$ .

376 We will argue now that

<sup>377</sup> 
$$\mathbb{P}\left[\pi\left[\frac{m}{\log n}+1\right]\cup M \text{ is not a matching } \wedge \pi\left[\frac{m}{\log n}+1\right]\notin M_{L}^{*} \mid Y_{1},\ldots,Y_{i-1}\right] \geq 1-\frac{1}{\log^{5} n}.$$
 (2)

Since  $E_1$  happens, observe that the second part of the stream consists of  $m(1 - \frac{1}{\log n}) - O(n \log^2 n)$  edges that cannot be added to matching M, at most n/2 edges of  $M_L^*$  (depending on the outcome of variables  $Y_1, \ldots, Y_{i-1}$ ), and at most  $O(n \log^2 n)$  edges that could extend M. Further, the arrival order of the edges  $\pi(\frac{m}{\log n}, m]$  in the second part of the stream is

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uniform random, since the computed matching M is not affected by their order. Hence,

$$\mathbb{P}\left[\pi[\frac{m}{\log n} + 1] \cup M \text{ is not a matching } \wedge \pi[\frac{m}{\log n} + 1] \notin M_L^* | Y_1, \dots, Y_{i-1} \right] \\ \ge \frac{m(1 - \frac{1}{\log n}) - O(n \log^2 n)}{m(1 - \frac{1}{\log n})} \ge 1 - \frac{1}{\log^5 n},$$

using the assumption that the graph has at least  $C \cdot n \log^{10} n$  edges, for a large enough C. 385 The key part of our argument is as follows: Let  $\Pi$  be the set of permutations that fulfill 386 the event in Inequality 2. Given  $\Pi$ , we generate a set of permutations  $\Pi'$  with  $\Pi' \supseteq \Pi$ , 387 which thus implies that the respective event is more likely to happen than the event in 388 Inequality 2. Let  $\pi \in \Pi$  be any permutation. Consider edge  $e_i$  and let  $j_i$  be such that 389  $\pi[j_i] \in M$  is the edge incident to  $e_i$ . Since  $e_i \in M_L^*$ , we know that  $t_i > j_i$ . Construct 390 now new permutations such that  $e_i$  is removed from its position  $t_i$  and is inserted at every 391 position  $\{t_i + 1, t_i + 2, ..., m\}$  and add the resulting permutations to  $\Pi'$ . Observe that for 392 any permutation  $\pi'$  created this way, the exact same matching M is computed, which uses 393 the fact that  $\pi[\frac{m}{\log n} + 1]$  cannot be added to M, which is important if  $e_i$  is inserted at a position larger than  $\frac{m}{\log n} + 1$ . Observe further that the conditionings  $Y_j$  stay the same, which uses the fact that  $\pi[\frac{m}{\log n} + 1] \notin M_L^*$ . Observe that  $\Pi'$  and  $\Pi$  are not identical, since we do not necessarily have that  $\pi'[\frac{m}{\log n} + 1] \cup M$  is not a matching for  $\pi' \in \Pi'$ . By construction, 394 395 396 397 at least a  $(1 - \frac{1}{\log n})$ -fraction of the permutations in  $\Pi'$  imply  $Y_i = 0$ . We thus obtain: 398

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$$\mathbb{P}[Y_i = 0 | Y_1, \dots, Y_{i-1}] \ge$$

$$(1 - \frac{1}{\log n}) \cdot \mathbb{P}\left[\pi\left[\frac{m}{\log n} + 1\right] \cup M \text{ is not a matching } \wedge \pi\left[\frac{m}{\log n} + 1\right] \notin M_L^* \mid Y_1, \dots, Y_{i-1}\right]$$
$$\geq (1 - \frac{1}{\log n})(1 - \frac{1}{\log^5 n}) \geq 1 - \frac{2}{\log n}.$$

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We now use the Chernoff bound for dependent variables stated in Theorem 2. Using  $k = (\frac{1}{2} - \delta)\mu(G)$ , we obtain (using  $\mu = 2k/\log n$ , and  $\delta = 1$  in Theorem 2):

$$\mathbb{P}\left[\sum_{i=1}^{k} Y_i \ge 2\frac{2k}{\log n}\right] \le \left(\frac{e}{4}\right)^{\frac{2k}{\log n}} \le n^{-10} ,$$

using the assumption  $\mu(G) = \Omega(n^{\frac{1}{4}})$ . The result follows.

In the remaining analysis, with the help of the previous lemma we bound the number of augmenting paths found in Lemma 15. We then conclude with our main theorem, where we show that one of the two computed matchings returned by the algorithm is necessarily large.

▶ Lemma 15. Let  $p = \Omega(1)$ , suppose that  $|M| + \mu(H) = (\frac{1}{2} + \delta)\mu(G)$ , and let  $|M| = \alpha\mu(G)$ . Then, with high probability,

$$|\mathcal{P}| \ge p\mu(G)\left(\frac{1-2\delta}{1+p} - \alpha\right) - o(\mu(G))$$
.

**Proof.** We follow the structure of the proof of Lemma 7. By an application of a Chernoff bound, we obtain  $|M'| = p|M| \pm O(\sqrt{|M|\ln(|M|)})$ , with probability at least  $1 - |M|^{-C}$ , for an arbitrarily large constant C. Next, by Theorem 6 and Lemma 14, with high probability in  $\mu(G)$  we have

$$\min\{|M_L|, |M_R|\} \ge \frac{p}{1+p}(\frac{1}{2}-\delta)\mu(G) - o(\mu(G)) .$$

Since we assumed that  $\mu(G) = \Omega(n^{\frac{1}{4}})$ , this event also holds with high probability in n. As argued in the proof of Lemma 7, the quantity  $|M_L| + |M_R| - |M'|$  bounds the number of 3-augmenting paths found, which then completes the proof:

$$\begin{aligned} & |\mathcal{P}| \geq |M_L| + |M_R| - |M'| \geq \frac{2p}{1+p} (\frac{1}{2} - \delta)\mu(G) - p|M| - o(\mu(G)) \\ & = p\mu(G) \left(\frac{2}{1+p} (\frac{1}{2} - \delta) - \alpha\right) - o(\mu(G)) = p\mu(G) \left(\frac{1-2\delta}{1+p} - \alpha\right) - o(\mu(G)) \;. \end{aligned}$$

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▶ **Theorem 16.** Setting  $p = \sqrt{2} - 1$  in Algorithm 2 gives a one-pass random order semistreaming algorithm for MBM with approximation ratio  $\frac{1}{2} + \frac{2\sqrt{2}-3}{4\sqrt{2}-10} \ge 0.5390$  that succeeds with high probability.

<sup>415</sup> **Proof.** Suppose that  $|M| = \alpha \mu(G)$  and  $|M| + \mu(H) = (\frac{1}{2} + \delta)\mu(G)$ . By Lemma 11, we have <sup>416</sup>  $\mu(H) \ge (1 - 2\alpha)\mu(G)$ . Hence,  $(1 - \alpha)\mu(G) \le (\frac{1}{2} + \delta)\mu(G)$ , which in turn implies  $\alpha \ge \frac{1}{2} - \delta$ . <sup>417</sup> Plugging this into the bound given in Lemma 15, we obtain (ignoring the  $o(\mu(G))$  term):

The quantity  $|M| + \max\{|\mathcal{P}|, \mu(H)\}$ , i.e., the size of the resulting matching, is minimizes if  $|\mathcal{P}| = \mu(H)$ . Hence, setting the right side of the previous inequality equal to  $(\frac{1}{2} + \delta)\mu(G)$ , we obtain  $\delta = \frac{p(p-1)}{2p^2 - 6p - 4}$ , which is maximized for  $p = \sqrt{2} - 1$  (observe that this is the same value as in the proof of Theorem 8). In this case, we obtain  $\delta = \frac{2\sqrt{2}-3}{4\sqrt{2}-10} \approx 0.03950$ , which completes the proof.

# 425 **6** Conclusion

In this paper, we gave a new method for finding a set of disjoint 3-augmenting paths that allows the augmentation of a maximal matching such that the resulting matching is of size at least  $\sqrt{2} - 2$  times the size of a maximum matching. Our method is simple and only requires running the GREEDY matching algorithm on a random subgraph. We applied this method in the data streaming setting and improved over the state-of-the-art one-pass random order algorithm and the state-of-the-art two- and three-pass adversarial order algorithms.

How large a matching can we compute in a single pass in the random order setting? All relevant known lower bounds for matchings [18, 22, 19] are highly sensitive to the arrival order of the edges and do not translate to the random order setting. Can we compute a 2/3-approximation in a single pass in the random order semi-streaming setting? In the adversarial order setting, it is known how to obtain a  $2/3 - \delta$  approximation in  $O(\frac{1}{\delta})$  passes. How many passes are required to obtain a 2/3-approximation?

<sup>438</sup> — References

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