Maximum Bipartite Matching Size And Application to Cuckoo Hashing*

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Abstract

Cuckoo hashing with a stash is a robust high-performance hashing scheme that can be used in many real-life applications. It complements cuckoo hashing by adding a small stash storing the elements that cannot fit into the main hash table due to collisions. However, the exact required size of the stash and the tradeoff between its size and the memory over-provisioning of the hash table are still unknown.

We settle this question by investigating the equivalent maximum matching size of a random bipartite graph, with a constant left-side vertex degree d = 2. Specifically, we provide an exact expression for the expected maximum matching size and show that its actual size is close to its mean, with high probability. This result relies on decomposing the bipartite graph into connected components, and then separately evaluating the distribution of the matching size in each of these components. In particular, we provide an exact expression for any finite bipartite graph size and also deduce asymptotic results as the number of vertices goes to infinity.

We also extend our analysis to cases where only part of the left-side vertices have a degree of 2; as well as to the case where the set of right-size vertices is partitioned into two subsets, and each left-side vertex has exactly one edge to each of these subsets. Finally, in the case where the constant left-side degree satisfies $d \ge 3$, we show how our method can be used to bound from above the expected maximum size matching.

Our results improve upon previous results in two ways. First, we give an exact expression of the expected size of the maximum matching and not only the threshold for achieving a perfect matching with high probability (namely, we show the trade-off between the size of the stash and the memory over-provisioning of the table, and not only the over-provisioning threshold beyond which the stash can be eliminated). Second, our results hold for any finite graph size and are not only asymptotic.

Keywords: maximum matching, random bipartite graph, hashing, power of two choices

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1 Introduction

A hash table is a classical randomized data structure that is often used to store efficiently large sets of elements in an associative manner, as it provides fast insertion, deletion and lookup of elements. Basically, a hash table consists of applying a hash function on each element (or more precisely, on each key) to determine where to store it. Hash schemes differ mainly in the way they solve conflicts. A seminal result [4] on *multiple-choice hashing schemes* shows that using a constant number d > 1 of hash functions instead of just one significantly boosts the performance. These multiple-choice hashing schemes can be modeled as solving a matching problem in a bipartite graph, with n elements as left-side vertices, m locations as right-side vertices, and d edges for each left-side vertex as hash values. In this setting, if each location can store at most one element, no hashing scheme can store more elements than the size of the maximum size matching.

A prime example of multiple-choice hashing scheme is *cuckoo hashing* [16, 34], which has recently drawn a lot of attention due to its efficient space utilization along with constant query and deletion times, as well as a constant expected insertion time (e.g., [3,15-18,20,21,32] and references therein). In cuckoo hashing, each of the *n* elements (or *balls*) uses *d* independent hash functions to pick up to *d* locations (or *bins*). Typically, each bin is of size one and d = 2. Upon insertion of a ball *x*, it is placed in one of its bins even if it is full. In such a case, it displaces another ball which is moved to the bin corresponding to its other choice. This process can cause another ball displacement and continues until either no ball is displaced or an infinite loop occurs (in the latter case, the scheme fails to insert the ball into the hash table).

It is important to notice that cuckoo hashing succeeds in inserting an element if and only if an augmenting path originating from the corresponding vertex exists [21]. This is because inserting an element into a cuckoo hash table is equivalent to finding an augmenting path in the corresponding graph (that is, a path that starts from the vertex corresponding to the considered element, and alternates between unmatched and matched edges until it ends at a right-side vertex whose all edges are unmatched). Notice that for any sub-path (r_1, v, r_2) , where v is a left-side vertex and r_1, r_2 are right-side vertices, (r_1, v) must be a matched edge and (v, r_2) an unmatched edge. Intuitively, this corresponds to moving element v from bin r_1 to bin r_2 . Since maximum size matching can be computed by finding such augmenting paths, when considering each left-side vertex only once and in arbitrary order, we can immediately conclude that the number of elements that a cuckoo hashing inserts successfully *is exactly the size of the maximum matching*. For example, all n elements can be inserted if and only if the corresponding graph has a *perfect matching* (namely, a maximum matching of size n; see [21] for more details).

Indeed, most papers so far have focused on the threshold conditions under which there is a perfect matching with high probability [15, 16, 18, 21], so that all elements could fit in the hash table. However, modern hashing schemes provide more robustness by considering an additional memory (often called *stash* or overflow list) that stores a small number of elements outside the main hash table $[23-27]^1$. Thus, sizing the stash incurs a trade-off between its size and the *load* m/n of the hash-table. It is an important design issue that is not yet understood [29, Open Question 5]. This open question is answered in this paper for d = 2 by evaluating the *expected maximum matching size*, and showing that the actual maximum matching size is sharply concentrated around its expected value. Thus, the difference between n and the expected maximum matching size provides the required size of the stash, which should store all elements with high probability. We also provide exact analysis when the average number of choices is less than 2 to minimize the number of memory accesses. We further obtain a lower bound on the required stash size when d > 2.

We note that for other multiple-choice hashing schemes, our results provide *a lower bound on the size of the stash*. This is because the maximum matching size of the graph is always an upper bound on

¹The stash can then be implemented using content-addressable memories or fast cache lines. See discussion in [25, 29]. In some cases, there is no stash and elements are simply dropped. This corresponds to *lossy* hash tables [17].

the number of elements that can be inserted into the hash table. Moreover, since finding the maximum matching in bipartite graphs is a fundamental problem with a wide range of applications in computer science, we believe that our results have also a theoretical significance and may be used in other contexts.

To prove our results on the expected size of the maximum bipartite match, we decompose each random bipartite graph into *connected components*, and then separately analyze each component and evaluate the size of its local maximum bipartite match. The size of the maximum bipartite matching is the sum of the sizes of all local matches. Then, we count the number of connected components in the graph and thus derive the size of the maximum matching in the entire graph. Our concentration results are based on applying Azuma's inequality [5] to a Doob martingale, which is defined over the maximum matching size when exposing vertices one at a time. Our results for d > 2 also use Huisimi tree enumerations [7,22].

1.1 Our Contributions

Consider a bipartite graph $G = \langle L + R, E \rangle$, with two disjoint vertex sets L and R of sizes |L| = n and |R| = m, and edge set E. Assume that each vertex $v \in L$ independently chooses d = 2 vertices in R. Furthermore, assume that repetitions are allowed, i.e. each vertex is picked uniformly at random in R, and therefore v might pick the same vertex twice, thus yielding two parallel edges.

First, given any n and any m, we present the expected size of the maximum bipartite matching. We obtain this result by decomposing the bipartite graph into connected components, and then separately evaluating the distribution of the matching size in each of these components. We also later show that with high probability the matching size is close to its expectation. The result is illustrated in Figure 2.

Theorem (appears as Theorem 3.6). Let d = 2 and $b = \min\{n, m-1\}$. The expected maximum matching size $\mu(G)$ is

$$\mu(G) = m - \sum_{s=0}^{b} \binom{n}{s} \cdot \binom{m}{s+1} \cdot \left(1 - \frac{s+1}{m}\right)^{2(n-s)} \cdot \left(\frac{s+1}{m}\right)^{2s} \cdot \frac{2^s \cdot s!}{(s+1)^{s+1}}$$

Next, we obtain the asymptotic expected size of the maximum bipartite matching as $n \to \infty$ with a constant load $\alpha = \frac{n}{m}$. It is illustrated in Figure 3.

Theorem (appears as Theorem 3.9). Let d = 2, the limit normalized expected maximum matching size $\gamma = \lim_{n \to \infty} \frac{\mu(G)}{n}$ is

$$\gamma = \lim_{n \to \infty} \frac{\mu(G)}{n} = \frac{1}{\alpha} + \frac{1}{2\alpha^2} \cdot W\left(-2\alpha \cdot e^{-2\alpha}\right) + \frac{1}{4\alpha^2} W^2\left(-2\alpha \cdot e^{-2\alpha}\right),\tag{1}$$

where the Lambert-W function is the inverse function of the function $\omega(x) = xe^x [12]^2$.

Equation (1) yields that a perfect matching requires the load to be at most $\frac{1}{2}$ (see Corollary 3.10), thus confirming known results on the feasibility of storing all elements in a cuckoo hashing scheme [16]. Beyond this threshold, the normalized expected size of the matching decreases as the load α increases. For $\alpha = 1$ (corresponding to n = m) it is approximately 84% (see Example B.2), implying that a stash of size at least $\approx 0.16n$ is required.

We extend the results above in two ways. First, we consider cases where part of the left-side vertices have degree 1 and the others have degree 2. These cases occur when the average number of choices is important, for instance to minimize the energy needed to insert all elements [23, 36]. In these cases, we present the exact and asymptotic expected sizes of the maximum bipartite matching.

Then, we assume that the set R of right-size vertices is partitioned into two subsets, and each leftside vertex has exactly one edge to each of these subsets. This corresponds to a common alternative

²See Appendix B.1 for a plot of the Lambert-W function in Figure 1.

model of cuckoo hashing [16,27,34]. We provide again an exact expected size of the maximum bipartite matching in the finite case. We then show that the asymptotic limit using this static partition is the same as without it. We further provide the exact conditions on the static partition for a perfect matching to exist, and in particular derive a result of [16] as a corollary.

Finally, we combine our decomposition into connected components with results on Husimi tree enumeration [7,22] to derive an upper bound on the expected maximum matching size for cases where there are more than 2 choices for each left-side vertex. This additional choice, already introduced and studied in previous works [15, 18, 21], enables higher matching sizes and therefore lower overflow fractions.

1.2 Related Work

Multiple-choice hashing schemes were first considered in the seminal paper of Azar et al. [4]. It showed that placing each element in the least occupied bin among a constant number d of random bins significantly improves the maximum bin load to $\frac{\log \log n}{\log d} + O(1)$ with high probability (compared to the case where d = 1, in which the maximum bin load is $\log n (1 + O(1))$). This result initiated an extensive research with many variants of multiple-choice hashing schemes, which typically exhibited the so-called *power of two choices* with d = 2 [30]. For brevity, we next survey only works that directly correspond to our paper.

First, we relate to works which considered the same model as in this paper (a random bipartite graph with constant left-side vertex degree). Motivated by achieving a performance guarantee for the cuckoo hashing scheme [34], the main effort has been to find a load threshold, such that for any load below the threshold a perfect matching exists with high probability. It is known that a cuckoo hashing scheme with d = 2 succeeds with high probability if the load is less than a load threshold of $\frac{1}{2}$, but fails when the load is larger than $\frac{1}{2}$ [16]. Recent works [15, 18, 21] have settled the problem of finding the corresponding thresholds for d > 2. Our paper differs in that we consider all possible load values for d = 2 (where we also confirm the known results for load less than 0.5). Moreover, while most of the works investigate only the asymptotic behavior, we also present in our paper analytical expressions for finite random graphs along with the asymptotic ones.

The problem of finding the expected maximum matching size is also investigated assuming other models of random graphs, mainly trees. In [9, 10] the authors investigate the expected maximum matching size of an (r, s)-tree, finding that for almost all (n, n)-trees the percentage of dark vertices in a maximum matching is at least 72%. A more recent work [6] presents results related to the expected maximum matching size of the class of simply-generated trees. A model of a loop graph is considered by [33], showing a lower bound on the expected maximum matching size. While using the cavity method of statistical physics [40], the authors find analytically the value under consideration for the Erdős graph G(n, c/(n-1)), where c < 2.7183. Our paper differs in that it considers a different model of random bipartite graphs, where each vertex in L chooses a constant number of vertices in R.

Additional related works deal with the probability of a perfect matching in other random graph models. For instance, in a random directed bipartite graph with n left-side and n right-side vertices, and an outward degree d at each vertex, the probability that the random bipartite graph contains a perfect matching approaches 1 if d > 1, but approaches 0 otherwise [13]. Also, in a random bipartite graph with n left-side vertices, n right-side vertices, cn edges picked uniformly at random, and a degree of at least 2, there is a perfect matching with high probability [19].

Finally, the conjectures in [11, 35] consider the expected minimum matching weight given a full bipartite graph with random exponentially distributed edge weights. These conjectures are proved in [1, 28, 31].

Paper Organization We start with preliminary definitions in Section 2. Section 3 provides the expected maximum matching size of random bipartite graphs with left-side vertex degree 2. Then, Sec-

tion 4 considers a variation of the problem in which each left-side vertex degree is at most 2. Next, in Section 5, we solve the problem in case the right-side vertices are partitioned into two subsets, and each left-side vertex has exactly one edge to each of these subsets. Last, Section 6 provides an upper bound on the expected maximum matching size when the constant left-side vertex degree is at least three.

Due to space limits, some proofs as well as further evaluation of the results in this paper are presented in Appendix A and Appendix B.

2 Definitions and Problem Statement

Given two disjoint sets of vertices L and R of size n and m respectively, we consider a random bipartite graph $G = \langle L + R, E \rangle$, where each vertex $v \in L$ has d = 2 outgoing edges whose destinations are chosen independently at random among all vertices in R. We allow both choices to be the same vertex, implying that G might have parallel edges. For brevity, we sometimes say that $v \in L$ chooses a vertex $v' \in R$ if (v, v') is in E. The *load* of G is denoted by $\alpha = \frac{n}{m}$.

We also consider cases when the averages number of choices, as defined below, is less than 2

Definition 2.1. Let d_v be the number of choices of each vertex $v \in L$. The average number of choices a is the average left-side vertex degree, i.e. $a = \frac{\mathbb{E}(\sum_{v \in L} d_v)}{n} = \frac{\sum_{v \in L} \mathbb{E}(d_v)}{n}$.

Specifically, we consider the following two extensions where, for each vertex $v \in L$, $d_v \leq 2$ and a < 2: (i) a random bipartite graph $G_a = \langle L + R, E \rangle$, where exactly d_2 of the vertices in L choose two vertices in R and the other $d_1 = n - d_2$ vertices in L choose exactly one vertex in R, such that $a = \frac{d_1 + 2 \cdot d_2}{n}$, and, (ii) a random bipartite graph $G_p = \langle L + R, E \rangle$, where each vertex in L chooses exactly two vertices in R with probability p, and one vertex with probability 1 - p. This implies that in G_p , the average number of choices a = 1 + p.

Finally, we also consider a *static partitioning of the choices*; the set R is partitioned into two disjoint sets R_u and R_d of sizes $\beta \cdot m$ and $(1 - \beta) m$. In that case, we consider a random bipartite graph $G_\beta = \langle L + (R_u \cup R_d), E \rangle$, where each vertex $v \in L$ chooses exactly one vertex in R_u and another vertex in R_d .

This paper focuses on the expected size of the maximum size matching of G, which is captured by the following definition:

Definition 2.2. The operator $\mu(\cdot)$ extracts the expected size of the maximum size matching. It operates both on deterministic and random bipartite graphs. (Namely, for a deterministic graph H, $\mu(H)$ is simply the size of the maximum size matching of H.)

Definition 2.3. The normalized limit expected maximum matching size $\gamma = \lim_{n \to \infty} \frac{\mu(\cdot)}{n}$ is the limit percentage of the expected maximum matching size (out of the number of the vertices in L).

Our goal is to find both the *expected maximum matching size* as well as the *normalized limit expected maximum matching size* for the above-mentioned graph models.

3 Bipartite Graphs with d = 2

Our approach relies on considering the connected components of the random bipartite graph G. Therefore, we start by stating some essential lemmas on these connected components, before establishing our result on the expected matching size.

3.1 Connected Components in Deterministic Graphs

The following lemmas are for a given bipartite graph $H = \langle L_H + R_H, E_H \rangle$, where each vertex in L_H has degree 2 (parallel edges are allowed), with $|L_H| = s$ and $|R_H| = q$. Their proofs are presented in Appendix A.

Lemma 3.1. If $s \leq q - 2$, then H is not connected.

Lemma 3.2. If H is connected and $s \ge q$, then $\mu(H) = q$.

Lemma 3.3. If H is connected and s = q - 1 then $\mu(H) = s$.

Lemma 3.4. For any graph with s = q - 1, H is connected if and only if it is a tree.

Lemma 3.5. The number T_s of connected bipartite graphs H whose $|L_H| = s$ and $|R_H| = s + 1$ is $T_s = (s+1)^{s-1} s!$

3.2 Expected Maximum Matching Size

In this section we deal with a random graph, in which each left-side vertex chooses d = 2 right-side vertices (parallel edges are allowed). Note that further evaluation of the results reported here appears in B.2.

The next theorem is the main result of this paper.

Theorem 3.6. Let d = 2 and $b = \min\{n, m-1\}$. The expected maximum matching size $\mu(G)$ is

$$\mu\left(G\right) = m - \sum_{s=0}^{b} \binom{n}{s} \cdot \binom{m}{s+1} \cdot \left(1 - \frac{s+1}{m}\right)^{2(n-s)} \cdot \left(\frac{s+1}{m}\right)^{2s} \cdot \frac{2^s \cdot s!}{\left(s+1\right)^{s+1}}.$$

Proof. Let M be a maximum matching of G. Our proof is based on counting the expected number of vertices in R that are not part of M, and on the decomposition of G into its connected components.

Lemma 3.1 yields that any connected component of G with s left-side vertices has at most s + 1 right-side vertices. We call a connected component with s left-side vertices and s + 1 right-side vertices a deficit component of size s. Lemma 3.3 implies that the maximum matching size of any such deficit component is s. Therefore, exactly one of its right-side vertices is not part of M. Notice that in all other connected components, where q < s + 1, the maximum matching size of G is exactly q (Lemma 3.2), implying that all their right-side vertices are part of M.

Thus, in order to calculate the size of M, it suffices to count the number of deficit components x. The size of M is m-x because exactly x right-side vertices do not participate in M, one for each deficit component.

Let $P_s = \frac{2^s T_s}{(s+1)^{2s}}$ be the probability that a bipartite graph $H = \langle L_H + R_H, E_H \rangle$ is connected, with degree 2 for all vertices in L_H , where $|L_H| = s$ and $|R_H| = s + 1$.

The expected number of deficit components of size s is $\binom{n}{s}\binom{m}{s+1} \cdot \left(1 - \frac{s+1}{m}\right)^{2(n-s)} \cdot \left(\frac{s+1}{m}\right)^{2s} \cdot P_s$. The above expression consists of the following factors (in order):

(*i*) choosing the s vertices in L;

(*ii*) choosing the s + 1 vertices in R;

(*iii*) the probability that all s + 1 vertices in R may be connected only to the chosen s vertices in L;

(*iv*) the probability that all s vertices in L are only connected to the s + 1 vertices in the right side; and, (v) the probability that all chosen vertices are connected.

Finally, we calculate x by summing over all possible values on s. As mentioned before, the expected size of M is given by m - x. We get: $\mu(G) = m - \sum_{s=0}^{b} {n \choose s} {m \choose s+1} \cdot \left(1 - \frac{s+1}{m}\right)^{2(n-s)} \cdot \left(\frac{s+1}{m}\right)^{2s} \cdot P_s$, where $b = \min\{n, m-1\}$, $P_s = \frac{2^s T_s}{(s+1)^{2s}}$, and $T_s = (s+1)^{s-2} \cdot (s+1)!$, as found in Lemma 3.5. \Box

3.3 Concentration Result

We next show that the size of the maximum matching is highly concentrated around its expectation $\mu(G)$. In order to prove this result, we apply Azuma's inequality to a Doob martingale (more specifically, the martingale is a vertex exposure martingale of the left-side vertices).

Note that as long as all left-side vertices pick their edges independently, this concentration result holds regardless of the value of d, and more generally regardless of the specific distribution over which the hash functions are defined. Therefore, the concentration result applies also for the settings of Sections 4–6.

Theorem 3.7. Let *H* be a specific instance of the random graph *G*, as defined in Section 2. For any $\lambda > 0$, $\Pr(|\mu(H) - \mu(G)| > \lambda\sqrt{n}) < 2e^{-\lambda^2/2}$.

Notice that if we are interested only in one-sided bounds, we can get a slightly tighter result: $\Pr(\mu(G) - \mu(H) > \lambda \sqrt{n}) < e^{-\lambda^2/2}$. This is exploited in the following corollary, which shows that to obtain a given overflow fraction, the needed stash size grows sub-linearly with *n* beyond its average value.

Corollary 3.8. To achieve an overflow fraction of ϵ in cuckoo hashing with stash, a stash of size $\left(n - \mu(G) + \sqrt{2n \cdot \ln(1/\epsilon)}\right)$ suffices.

Proof. If a stash of size $n - \mu(G) + \sqrt{2n \cdot \ln 1/\epsilon}$ is used, cuckoo hashing fails if and only if $n - \mu(H) > n - \mu(G) + \sqrt{2n \cdot \ln 1/\epsilon}$, or by rewriting it, $\mu(G) - \mu(H) > \sqrt{2n \cdot \ln 1/\epsilon}$. By substituting $\lambda = \sqrt{2 \cdot \ln 1/\epsilon}$ in the above one-sided bound, we get the claimed result.

3.4 Limit Normalized Expected Maximum Matching Size

We are now interested in the asymptotic expression where $n \to \infty$ with $\alpha = \frac{n}{m}$ constant. The following results show an interesting connection between the limit normalized expected maximum matching size and the Lambert-W function, and even a connection between the perfect matching threshold and the radius of convergence of the Lambert-W function [12]. For further details on the Lambert-W function, see also Appendix B.1.

Theorem 3.9. Let d = 2. The limit normalized expected maximum matching size $\gamma = \lim_{n \to \infty} \frac{\mu(G)}{n}$ is given by:

$$\gamma = \frac{1}{\alpha} + \frac{1}{2\alpha^2} \cdot W\left(-2\alpha \cdot e^{-2\alpha}\right) + \frac{1}{4\alpha^2} W^2\left(-2\alpha \cdot e^{-2\alpha}\right),\tag{2}$$

where the Lambert-W function is the inverse function of the function $\omega(x) = xe^x$.

The following corollary shows that for $\alpha = \frac{n}{m} \leq \frac{1}{2}$, the probability for a right-side vertex to be part of a maximum matching goes to 1. This corollary also follows from the previously known result that there is a perfect matching with high probability in cuckoo hash tables with load $\alpha \leq \frac{1}{2}$ [16].

Corollary 3.10. Let d = 2 and $\alpha = \frac{n}{m} \leq \frac{1}{2}$. Then the limit normalized expected maximum matching size is $\gamma = \lim_{m \to \infty} \frac{\mu(G)}{n} = 1$.

Proof. In case
$$\alpha \leq \frac{1}{2}$$
, $W\left(-2\alpha \cdot e^{-2\alpha}\right)$ equals -2α , thus, $\gamma = \frac{1}{\alpha} + \frac{1}{2\alpha^2} \cdot (-2\alpha) + \frac{1}{4\alpha^2} (-2\alpha)^2 = 1$

4 Bipartite Graphs With $d_v \leq 2$

In this section we relax the constraint that each vertex in L chooses exactly 2 vertices in R, and let each left-side vertex choose either 1 or 2 right-side vertices. Since we can divide the set of vertices either deterministically or randomly, we will discuss the results in both cases. These results correspond for example to cases in which the average number of choices is important (e.g. [23]). See also [36] for a similar model.

First, in the deterministic case, we find the expected maximum matching size of the graph $G_a = \langle L + R, E \rangle$, where each vertex $v \in L$ independently chooses a predetermined number $d_v \in \{1, 2\}$ of random vertices in R, such that $a = \frac{d_1 + 2 \cdot d_2}{n}$.

Second, in the random case, we analyze the slightly different case of a random bipartite graph $G_p = \langle L + R, E \rangle$ where each vertex chooses two vertices with probability p and one vertex with probability 1 - p.

Note that further evaluation of the results reported in this section can be found in Appendix B.3.

4.1 Connected Components in Deterministic Graphs

As in Section 3.1, we now consider a deterministic bipartite graph $H = \langle L_H + R_H, E_H \rangle$, with $|L_H| = s$ and $|R_H| = q$. We assume that the degree of each vertex in L_H is at most 2.

Observation 4.1. Lammas 3.1, 3.2, and 3.3 hold also when the degree of each vertex in L_H is at most (but not necessarily) 2. Note that the proofs remain almost identical to the original proofs, replacing a few equalities with the corresponding inequalities.

Lemma 4.2. Let s + 1 = q. If H is connected then the degree of each vertex in L_H is 2.

Proof. Assume on the contrary that H is connected but that there is (at least) a single vertex $v_L \in L_H$ with degree 1. Consider the bipartite graph $\hat{H} = \langle \hat{L}_H + \hat{R}_H, \hat{E}_H \rangle$, that is given by removing the vertex v_L (and its connected edge) from H. By the construction of \hat{H} , we get that \hat{H} is connected, but $|\hat{L}_H| + 1 < |\hat{R}_H|$, which contradicts Lemma 3.1.

4.2 Expected Maximum Matching Size

4.2.1 Predetermined Number of Choices

In this section, we assume that each vertex $v \in L$ independently chooses $1 \le d_v \le 2$ random vertices in R, where d_v is predetermined. The following result provides the expected maximum matching size in this case.

Theorem 4.3. Given a predetermined average number of choices a, let $d_1 = (2 - a) \cdot n$ and $d_2 = n - d_1 = (a - 1) \cdot n$ be the number of vertices in L that choose one and two vertices in R, respectively. The expected maximum matching size $\mu(G_a)$ is given by:

$$\mu(G_a) = m - \sum_{s=0}^{b} {\binom{d_2}{s}} {\binom{m}{s+1}} \cdot \left(1 - \frac{s+1}{m}\right)^{2(d_2-s)+d_1} \cdot \left(\frac{s+1}{m}\right)^{2s} \cdot \frac{2^s \cdot s!}{(s+1)^{s+1}},$$

where $b = \min \{d_2, m - 1\}$.

4.2.2 Random Number of Choices

In this section, we assume that each vertex $v \in L$ independently chooses $1 \le d_v \le 2$ random vertices in R, where for each $v \in L$, d_v equals 2 with probability p, and it equals 1 with probability 1 - p. Based on Theorem 4.3, the following result reflects the expected maximum matching size in this case.

Theorem 4.4. The expected maximum matching size $\mu(G_p)$ is given by $\mu(G_p) = \sum_{d_2=0}^n {n \choose d_2} \cdot p^{d_2} \cdot (1-p)^{n-d_2} \cdot \mu\left(G_{a=1+\frac{d_2}{2}}\right)$, where $\mu(G_a)$ is given by Theorem 4.3.

Proof. The number of vertices in L with degree 2 follows a Binomial distribution with n experiments and a probability of success p. In Theorem 4.3 we found the expected maximum matching size of each such instance. Thus, by the law of total expectation, the claimed result is given by computing the weighted average, where we compute a by the equations $d_1 + d_2 = n$ and $d_1 + 2 \cdot d_2 = a \cdot n$.

4.3 Limit Normalized Expected Maximum Matching Size

4.3.1 Predetermined Number of Choices

We are also interested in the asymptotic expression, where $n \to \infty$, such that we fix both the load $\alpha = \frac{n}{m}$ and the average number of choices $a = \frac{d_1+2\cdot d_2}{n}$ of the vertices. This is reflected in the following theorem.

Theorem 4.5. The limit normalized expected maximum matching size $\gamma_a = \lim_{n \to \infty} \frac{\mu(G_a)}{n}$ with average number of choices $a \in (1,2]$ is given by: $\gamma_a = \lim_{n \to \infty} \frac{\mu(G_a)}{n} = \frac{1}{\alpha} + \frac{W(-2\alpha(a-1) \cdot e^{-a\alpha})}{2\alpha^2 \cdot (a-1)} + \frac{W^2(-2\alpha(a-1) \cdot e^{-a\alpha})}{4\alpha^2 \cdot (a-1)}$. For a = 1, it is given by $\gamma_a = \lim_{n \to \infty} \frac{\mu(G_a)}{n} = \frac{1}{\alpha} - \frac{1}{\alpha} \cdot e^{-\alpha}$.

Interestingly, if even a small fraction of the elements do not have choice then the expected maximum matching size is not 1. This is reflected in the following corollary.

Corollary 4.6 ((No) Perfect Matching). If $1 \le a < 2$ then $\gamma_a < 1$.

4.3.2 Random Number of Choices

We now consider the case of the random bipartite graph $G_p = \langle L + R, E \rangle$, where each vertex chooses two vertices with probability p (and one vertex with probability 1-p). As we show in the next theorem, the asymptotic expression can be derived by γ_a .

Theorem 4.7. The limit expected maximum matching size $\gamma_p = \lim_{n \to \infty} \frac{\mu(G_p)}{n}$ where each vertex chooses two vertices with probability p (and one vertex with probability 1-p) is $\gamma_p = \gamma_{a=1+p}$.

5 Static Partitioning of the Choices

We now consider the random bipartite graph $G_{\beta} = \langle L + (R_u \cup R_d), E \rangle$, where R is now partitioned into two disjoint subsets R_u and R_d with $|R_u| = \beta \cdot m$ and $|R_d| = (1 - \beta) m$. Each vertex $v \in L$ independently chooses a single random vertex in R_u and another single random vertex in R_d . This corresponds, for example, to a hashing scheme that selects non-overlapping sets of bins as images of its hash functions (e.g., as in multilevel hashing scheme [8] or d-left [38]).

Note that further evaluation of the results reported in this section can be found in Appendix B.4.

5.1 Connected Components in Deterministic Graphs

The following lemma counts all the possible bipartite graphs H_{ud} of the form $\langle L_H + (R_{H_u} \cup R_{H_d}), E_H \rangle$ with degree 2 for each vertex in L_H , where $|L_H| = s$, $|R_{H_u}| = i$ and $|R_{H_d}| = j$, such that each vertex $v \in L_H$ is connected using a single edge to some vertex in R_{H_u} and another single edge to some vertex in R_{H_d} . **Observation 5.1.** Lemmas 3.1, 3.2, 3.3, and 3.4 hold for this case as well.

Lemma 5.2. Let s = i + j - 1. The number $T_{i,j}$ of connected bipartite graphs is $T_{ij} = i^{j-1} \cdot j^{i-1} \cdot s! = i^{j-1} \cdot j^{i-1} \cdot (i+j-1)!$

Proof. The proof is identical to the proof of Lemma 3.4 with two modifications. First, instead of initially counting the number of trees over the set R_H , we count the number of parity trees [37] over the disjoint sets R_{H_u} and R_{H_d} . By [37] we are given that the number of parity trees is $i^{j-1} \cdot j^{i-1}$. Second, we do not have to color the edges because of the partition.

5.2 Expected Maximum Matching Size

In the next theorem we find the expected maximum matching size with a static partition of the right-side vertices.

Theorem 5.3. Given the static partitioning of the bipartite graph G_{β} , the expected maximum matching size $\mu(G_{\beta})$ is

$$\mu(G_{\beta}) = m - \sum_{s=0}^{n} \binom{n}{s} \cdot \sum_{i=b_{1}}^{b_{2}} \binom{\beta \cdot m}{i} \cdot \binom{(1-\beta) \cdot m}{s+1-i} \cdot \left(1 - \frac{i}{\beta \cdot m}\right)^{n-s} \cdot \left(1 - \frac{s+1-i}{(1-\beta) \cdot m}\right)^{n-s} \cdot \left(\frac{i}{\beta \cdot m}\right)^{s} \cdot \left(\frac{s+1-i}{(1-\beta) \cdot m}\right)^{s} \cdot P_{i,s+1-i},$$

where $b_1 = \max\{0, s+1 - (1-\beta) \cdot m\}$, $b_2 = \min\{s+1, \beta \cdot m, \}$, $P_{ij} = \frac{T_{ij}}{(i \cdot j)^{i+j-1}}$, and $T_{ij} = i^{j-1} \cdot j^{i-1} \cdot (i+j-1)!$ (as given in Lemma 5.2).

5.3 Limit Normalized Expected Maximum Matching Size

As in the last sections, we are also interested in the asymptotic expression where $n \to \infty$ with both fixed $\alpha = \frac{n}{m}$ and fixed β . This is achieved in the following theorem.

Theorem 5.4. Given the static partitioning of the bipartite graph G_{β} , the limit normalized expected maximum matching size $\gamma_{\beta} = \lim_{n \to \infty} \frac{\mu(G_{\beta})}{n}$ for $\beta \in (0,1)$ is given by: $\gamma_{\beta} = \frac{1}{\alpha} - \frac{\beta \cdot (1-\beta)}{\alpha^2} \cdot (t_1 + t_2 - t_1 \cdot t_2)$, where t_1 , t_2 are provided by the following equations

$$\frac{\alpha}{1-\beta} \cdot e^{-\frac{\alpha}{\beta}} = t_1 \cdot e^{-t_2} \quad , \quad \frac{\alpha}{\beta} \cdot e^{-\frac{\alpha}{1-\beta}} = t_2 \cdot e^{-t_1} \tag{3}$$

and satisfy the condition $t_1 \cdot t_2 \leq 1$.

For $\beta \in \{0, 1\}$, (namely, the trivial partitions), the limit normalized expected maximum matching size γ_{β} is $\frac{1}{\alpha} - \frac{1}{\alpha} \cdot e^{-\alpha}$.

We deduce the following two corollaries.

Corollary 5.5 (Asymptotic Equivalence). Let d = 2. The limit normalized expected maximum matching size of G_{β} with $\beta = 0.5$ is the same as the limit expected maximum matching size of G.

Proof. We substitute $\beta = 0.5$ in the expression from Theorem 5.4, and get $\frac{\alpha}{0.5} \cdot e^{-\frac{\alpha}{0.5}} = t_1 \cdot e^{-t_2}$, $\frac{\alpha}{0.5} \cdot e^{-\frac{\alpha}{0.5}} = t_2 \cdot e^{-t_1}$. One of the solutions of the above equations is $t_1 = t_2 = -W(-2\alpha e^{-2\alpha})$. In the proof of Theorem 3.9, we showed that $-W(-2\alpha e^{-2\alpha}) \leq 1$. Thus, $t_1 \cdot t_2 < 1$. By substituting this solution in the expression for γ_β from Theorem 5.4, we get the exact expression as in Equation (2). \Box

Corollary 5.6. Let d = 2, $\alpha \leq \frac{1}{2}$, and fix a partition β . The limit normalized expected maximum matching size $\gamma_{\beta} = \lim_{n \to \infty} \frac{\mu(G_{\beta})}{n}$ is 1 whenever $\frac{1-\sqrt{1-4\alpha^2}}{2} \leq \beta \leq \frac{1+\sqrt{1-4\alpha^2}}{2}$.

Proof. One of the solutions to Equation (3) is given by: $t_1 = \frac{\alpha}{1-\beta}$, $t_2 = \frac{\alpha}{\beta}$. By substituting t_1 and t_2 in the expression for γ_{β} from Theorem 5.4, we get that the limit normalized expected maximum matching size is 1. We also have to verify that $t_1 \cdot t_2 \leq 1$. Since $\frac{\alpha}{1-\beta}$ and $\frac{\alpha}{\beta}$ are both positive, we are left with $\frac{\alpha}{1-\beta} \cdot \frac{\alpha}{\beta} < 1$. By solving the quadratic inequality, we get the claimed condition. Note that for $\alpha = 1/2$ the range reduces to $\beta = 1/2$.

6 Bipartite Graphs with d > 2

In this section we briefly show how our method can be applied to find an upper bound on the expected maximum size matching where each left-side vertex has d > 2 choices. Formally, we are given two disjoint sets of vertices L and R of size n and m, respectively, and a random bipartite graph $G^d = \langle L + R, E \rangle$, where each vertex $v \in L$ has d outgoing edges whose destinations are chosen independently at random (with repetition) among all vertices in R. We obtain the following upper bound on the maximum matching size of the bipartite graph G^d .

Theorem 6.1. Let $b = \min\left\{n, \left\lfloor \frac{m-1}{d-1} \right\rfloor\right\}$ and $q = (d-1) \cdot s + 1$. Then,

$$\mu\left(G^d\right) \le \min\left\{n \quad , \quad m - \sum_{s=0}^b \left(q - s\right) \binom{n}{s} \cdot \binom{m}{q} \cdot \left(1 - \frac{q}{m}\right)^{d(n-s)} \cdot \left(\frac{q}{m}\right)^{d \cdot s} \cdot \frac{d^s \cdot q!}{q^{(d-1) \cdot s + 2}}\right\}.$$

Appendix B.5 presents an evaluation of the upper bound and a comparison to the simulated expected matching size.

7 Concluding Remarks and Open Problems

In this work, we provided an exact expression for the expected maximum matching size of a random bipartite graph with each left-side vertex picking d = 2 right-side vertices. This result holds for any given finite numbers of left-side and right-side vertices. Then, we deduced asymptotic results as the number of vertices goes to infinity, and showed a connection to the Lambert-W function.

Both these results directly apply as exact results for the average number of inserted elements using cuckoo hashing. They also serve as an upper bound for any alternative hashing algorithm.

We also discussed alternative cases, using either a different number of vertex choices or a static partitioning of the right-side vertices. Finally, we showed how our technique can be used to bound from above the expected maximum matching size in case d > 2.

We are currently interested in the following two open problems: (*i*) *The average maximum matching size for any arbitrary d, and more generally any vertex number distribution*. While we have been able to obtain close bounds using the technique provided in this paper, we have not found general exact results, and, (*ii*) *The connection to the Lambert-W function*. This connection is intriguing, already appears e.g. in trees [12], and it might yield many keys to a more general counting approach in graph theory.

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Appendix

A Omitted proofs

A.1 Proof of Lemma 3.1

The proof follows by induction on s. For s = 1, there are 2 edges in the graph and therefore every graph with $q \ge 3$ is not connected. Assume that the claim holds up until s = s', we next prove that it holds for any bipartite graph H' such that $|L_{H'}| = s' + 1$ and $|R_{H'}| \ge s' + 3$. Assume towards a contradiction that there is a graph H' that is connected. We first show that there is a vertex in $R_{H'}$ with a degree 1: This follows from the fact that the average right-side degree is $\frac{2(s'+1)}{s'+3} < 2$, implying that there is at least one vertex with degree strictly less than 2; since the graph is connected, there are no right-side vertices with degree 0. Let v_r be such a vertex and let $v_\ell \in L_{H'}$ be the (only) left-side vertex to which it is connected. By the induction hypothesis, the graph induced by $L_{H'} \setminus \{v_\ell\}$ and $R_{H'} \setminus \{v_r\}$ is not connected, implying it has at least two connected components. In H', v_ℓ is connected to v_r and since its degree is 2 it can be connected only to one of these components. This implies that H' is also not connected, and the claim follows.

A.2 Proof of Lemma 3.2

We first consider the case where s = q. For $S \subseteq L_H$, let $d(S) \subseteq R_H$ be the set of vertices that are adjacent to any vertex in S. Hall's Theorem [14] implies that to prove that $\mu(H) = q$ (namely, there is a perfect matching in H) it suffices to prove that for every $S \subseteq L_H$, $|S| \leq |d(S)|$. Assume towards a contradiction that there is a subset $S \subseteq L_H$ such that |S| > |d(S)|, and denote |d(S)|by b. Furthermore, consider the bipartite graph $\hat{H} = \langle \hat{L}_H + \hat{R}_H, \hat{E}_H \rangle$, in which $\hat{L}_H = L_H \setminus S$, $\hat{R}_H = R_H \cup \{\hat{v}_R\} \setminus d(S)$ (where \hat{v}_R is a newly-introduced vertex) and any edge in E(H) of the form (v_ℓ, v_r) such that $v_\ell \in L_H \setminus S$ and $v_r \in d(S)$ is replaced with the edge (v_ℓ, \hat{v}_R) in \hat{E}_H . Notice that since H is connected, \hat{H} must be connected as well. Recall that |S| > b, thus $|\hat{L}_H| = |L_H \setminus S| \leq s - b - 1$, while $|\hat{R}_H| = |R_H \cup \{\hat{v}_R\} \setminus d(S)| = |R_H| - |d(S)| + 1 = s - b + 1$. This contradicts Lemma 3.1, implying that for every $S \subseteq L_H$, $|S| \leq |d(S)|$ and by Hall's Theorem $\mu(H) = q$.

For s > q, trivially $\mu(H) \leq q$. Therefore, it suffices to show that there exists a subset $S \subseteq L_H$ of size q, such that the corresponding bipartite subgraph is connected (and hence has a perfect matching of size q). We construct S in q iterations such that at the end of iteration n we end up with some subsets $S_n \subseteq L_H$ and $Q_n \subseteq R_H$ of the same size n, whose corresponding subgraph is connected. We start by n = 1 and pick some vertex $v_R \in R_H$ and one of its adjacent vertices $v_L \in L_H$. Assuming that at the end of iteration n, sets S_n and Q_n were chosen (and their corresponding graph is connected), we next construct S_{n+1} and Q_{n+1} . Let v_1 be an arbitrary vertex in S_n and let v_2 be an arbitrary vertex in $L_H \cup S_n$ (such a vertex always exists since s > q > n). Similarly, let v'_1 be an arbitrary vertex in Q_n and let v'_2 be an arbitrary vertex in $R_H \cup Q_n$. Since H is connected there is a path between v_1 and v_2 , and let v be the first vertex along this path that is not in S_n . Similarly, v' is the first vertex along the path between v'_1 and v'_2 that is not in Q_n . We differentiate between three cases: (i) v is adjacent to Q_n and v' is to S_n . In this case $S_{n+1} = S_n \cup \{v\}$ and $Q_{n+1} = Q_n \cup \{v'\}$ and the corresponding subgraph is connected; (ii) v is not adjacent to a Q_n . Let w be the vertex before v in the path between v_1 and v_2 , and let w' be the vertex before w in the path. Note that $w' \in S_n$ by the choice of v, and that $w \notin Q_n$ (otherwise v is adjacent to a Q_n). Thus, for $S_{n+1} = S_n \cup \{v\}$ and $Q_{n+1} = Q_n \cup \{w\}$, the corresponding subgraph is connected; (iii) v' is not adjacent to a S_n . The claim holds similarly to case (ii) by looking at the path between v'_1 and v'_2 . We continue this construction for q iterations, resulting in two subsets $S_q \subseteq L_H$ and $Q_q \subseteq R_H$ of size q each, whose corresponding subgraph is connected.

A.3 Proof of Lemma 3.3

Since each vertex in L_H has a degree of two, the sum of the degrees of all the vertices in R_H is 2s = 2q-2. Therefore, there must be at least one vertex $v_r \in R_H$ with degree 1 (there cannot be a vertex with degree 0 since H is connected). Let $v_L \in L_H$ be the (only) vertex that is connected to v_R and $\hat{v}_R \in R_H$ be the other vertex that is connected to v_L . Also consider the bipartite graph $\hat{H} = \langle \hat{L}_H + \hat{R}_H, \hat{E}_H \rangle$ that is given by removing v_R from H and adding a new edge (v_L, \hat{v}_R) . By the construction of \hat{H} , the degree of each vertex in \hat{L}_H is exactly 2. Moreover, since H is connected, \hat{H} is also connected. Hence, Lemma 3.2 implies that there is a matching of size s in \hat{H} . By the construction of \hat{H} , this is also a matching in graph H.

A.4 Proof of Lemma 3.4

First, if H is a tree then it is connected by definition. To show the other direction, we assume towards a contradiction that H is a connected graph with cycles; let C be a cycle in H, and consider an edge $e = (v_L, v_R)$ that resides at cycle C (where $v_L \in L_H$ and $v_R \in R_H$). We build the bipartite graph $\hat{H} = \langle \hat{L}_H + \hat{R}_H, \hat{E}_H \rangle$, such that $\hat{L}_H = L_H, \hat{R}_H = R_H \cup \{\hat{v}_R\}$, where \hat{v}_R is a newly-introduced vertex, and $\hat{E}_H = E_H \setminus \{e\} \cup \{\hat{e}\}$, where $\hat{e} = (v_L, \hat{v}_R)$. Intuitively, we replace one of the edges in the cycle to reach for a newly-introduced vertex, and by that we increase the size of the connected component. Notice that \hat{H} is connected and all vertices in \hat{L}_H have a degree of 2. But, $|\hat{L}_H| < |\hat{R}_H| - 1$, thus contradicting Lemma 3.1 and the claim follows.

A.5 Proof of Lemma 3.5

We count the connected bipartite graphs with two disjoint sets L_H and R_H . By Lemma 3.4, we have to count the number of trees over the set $L_H \cup R_H$, where edges must be of the form (v_L, v_R) , such that $v_L \in L_H$ and $v_R \in R_H$. We build (and count) the set as follows: The number of trees over the set R_H is $(s+1)^{s-1}$. For each such tree instance, we put a new vertex (originally from L_H) between each pair of adjacent vertices. There are s! possibilities to do so.

A.6 Proof of Theorem 3.7

Our notations follow those of [2]. We first define an exposure martingale, which exposes one left-side vertex at a time, along with all its outgoing edges. This martingale is equivalent to a regular vertex exposure martingale, in which all right-side vertices are exposed first, and then left-side vertices are exposed one by one.

Specifically, let G be the probability space of all two-choice bipartite graphs as defined in Section 2 and f the size of the maximum size matching of a specific instance. Assume an arbitrary order of the left-side vertices $L = \{v_1, \ldots, v_n\}$, and define X_0, \ldots, X_n by

$$X_i(H) = E[f(G) \mid \text{for } x \leq i \text{ and any } v_y \in R, (v_x, v_y) \in G \text{ iff } (v_x, v_y) \in H].$$

Note that $X_0(H) = \mu(G)$ since no edges were exposed, while $X_n(H) = \mu(H)$ as all edges are exposed.

Clearly, f satisfies the vertex Lipschitz condition since if two graphs H and H' differ at only one leftside vertex, $|f(H) - f(H')| \le 1$ (either that vertex is in the maximum matching or not). Thus, since each left-side vertex makes independent choices, [2, Theorem 7.2.3] implies that the corresponding vertex exposure martingale satisfies $|X_{i+1} - X_i| \le 1$. Hence, by applying Azuma's inequality, we immediately get the concentration result.

A.7 Proof of Theorem 3.9

We compute the limit of $\frac{\mu(G)}{n}$ as $n \to \infty$ such that $\alpha = \frac{n}{m}$:

$$\gamma = \lim_{n \to \infty} \frac{1}{n} \left(m - \sum_{s=0}^{b} \binom{n}{s} \binom{m}{s+1} \cdot \left(1 - \frac{s+1}{m} \right)^{2(n-s)} \cdot \left(\frac{s+1}{m} \right)^{2s} \cdot P_s \right)$$

We find through differentiation that $\left(1 - \frac{s+1}{m}\right)^{2(n-s)}$ is an increasing function with respect to n (where $m = \frac{n}{\alpha}$). Moreover, the expansion of $\frac{1}{n} \cdot {\binom{n}{s}} {\binom{m}{s+1}} \cdot {\binom{s+1}{m}}^{2s}$ shows that it is also an increasing function. Therefore, their product is also increasing and, by the monotone convergence theorem [39], we get

$$\gamma = \frac{m}{n} - \sum_{s=0}^{b} \lim_{n \to \infty} \left(\frac{1}{n} \cdot \binom{n}{s} \binom{m}{s+1} \cdot \left(1 - \frac{s+1}{m} \right)^{2(n-s)} \cdot \left(\frac{s+1}{m} \right)^{2s} \cdot P_s \right)$$

By substituting the expression for P_s , and using the facts that $\binom{n}{s} = \frac{n^s}{s!} + O(n^{s-1})$ and $\lim_{n\to\infty} (1+a/n)^n =$ e^a , we deduce:

$$\gamma = \frac{m}{n} - \frac{1}{n} \sum_{s=0}^{\infty} \frac{n^s}{s!} \cdot \frac{m^{s+1}}{(s+1)!} \cdot e^{-2\alpha(s+1)} \frac{(s+1)^{2s}}{m^{2s}} \cdot \frac{2^s \cdot (s+1)^{s-1} \cdot s!}{(s+1)^{2s}}$$

By substituting $m = \frac{n}{\alpha}$, and simplifying the above expression, we get:

$$\gamma = \frac{1}{\alpha} - \frac{1}{\alpha} \cdot \sum_{s=0}^{\infty} \alpha^s \cdot 2^s \cdot \frac{(s+1)^{s-1}}{(s+1)!} \cdot e^{-2\alpha(s+1)} = \frac{1}{\alpha} - \frac{1}{2\alpha^2} \cdot \sum_{j=1}^{\infty} \left(-2\alpha \cdot e^{-2\alpha}\right)^j \cdot \frac{(-j)^{j-2}}{j!}$$

Let $T(x) = \sum_{j=1}^{\infty} \frac{(-j)^{j-2}}{j!} \cdot x^j$ be a formal power series, where by substituting $x = -2\alpha \cdot e^{-2\alpha}$ we get the above expression. By differentiating T(x) and multiplying by x, we get:

$$x \cdot \frac{d}{dx}T(x) = -\sum_{j=1}^{\infty} \frac{(-j)^{j-1}}{j!} \cdot x^j = -W(x),$$

where the Lambert-W function is the inverse function of the function $\omega(x) = xe^x$ [12], and the last equality follows from its known Taylor expansion that converges as long as x is within the radius of convergence with $|x| \le e^{-1}$ [12]. Given that $x \cdot \frac{d}{dx}T(x) = -W(x)$, we compute T(x):

$$T(x) = \int \frac{1}{x} \cdot (-W(x)) \, dx = -W(x) - \frac{1}{2}W^2(x) \, ,$$

with convergence within $|x| < e^{-1}$.

Interestingly, the function $f(\alpha) = -2\alpha \cdot e^{-2\alpha}$ gets its minimum at $\alpha = 0.5$, where it precisely equals the radius of convergence $-e^{-1}$. Therefore, for all α we can substitute $x = -2\alpha \cdot e^{-2\alpha}$, since we are within the radius of convergence of T(x), and we finally derive the result.

Proof of Theorem 4.3 **A.8**

As in the proof of Theorem 3.6, our proof is based on counting the expected number of vertices in L that are not in some specific maximum matching M of G, based on the decomposition of G into its connected components. The proof is almost identical, with the modification that, due to Lemma 4.2, we only take into account the d_2 vertices that have a degree of 2 (instead of all n vertices in the proof of Theorem 3.6).

Thus, the expected number of connected components in G with s elements in L and s + 1 in R is given by:

$$\binom{d_2}{s}\binom{m}{s+1}\cdot\left(1-\frac{s+1}{m}\right)^{2(d_2-s)+d_1}\cdot\left(\frac{s+1}{m}\right)^{2s}\cdot P_s,$$

where the above expression consists of the same considerations as in the proof of Theorem 3.6. Finally, as before, adding the expressions for all possible s's and subtracting the sum from m yields the claimed result.

A.9 Proof of Theorem 4.5

We compute the limit of $\frac{\mu(G_a)}{n}$ as $n \to \infty$. We consider the case where $\alpha = \frac{n}{m}$ and $a = \frac{d_1 + 2 \cdot d_2}{n} > 1$ are fixed.

$$\gamma_a = \lim_{n \to \infty} \frac{\mu(G_a)}{n} = \lim_{n \to \infty} \frac{1}{n} \left(m - \sum_{s=0}^b \binom{d_2}{s} \binom{m}{s+1} \cdot \left(1 - \frac{s+1}{m}\right)^{2(d_2-s)+d_1} \cdot \left(\frac{s+1}{m}\right)^{2s} \cdot P_s \right)$$

Given that $a = \frac{d_1+2\cdot d_2}{n}$ and $n = d_1 + d_2$, we find that $d_2 = (a-1) \cdot n$ and $d_1 = (2-a) \cdot n$. Similarly to the proof of Theorem 3.9, we first have to find that each term in the summation is an increasing function with respect to n. We discover that $\left(1 - \frac{s+1}{m}\right)^{2(d_2-s)+d_1} = \left(1 - \frac{s+1}{m}\right)^{a\cdot n-s}$ is an increasing function (using differentiation), and also find that $\frac{1}{n} \cdot \binom{(a-1)\cdot n}{s} \binom{m}{s+1} \cdot \binom{s+1}{m}^{2s}$ is an increasing function as previously. Consequentially, each term in the sum is an increasing function and, by the monotone convergence theorem [39], we can put the limit inside the sum. By further simplifying the above expression as in the proof of Theorem 3.9 we eventually get:

$$\gamma_a = \frac{1}{\alpha} - \frac{1}{2\alpha^2 \cdot (a-1)} \cdot \sum_{j=1}^{\infty} \frac{(-j)^{j-2}}{j!} \cdot (-\alpha \cdot 2 \cdot (a-1) \cdot e^{-a\alpha})^j$$

Let $T(x) = \sum_{j=1}^{\infty} \frac{(-j)^{j-2}}{j!} \cdot x^j$ be a Taylor expansion, where by substituting $x = -\alpha \cdot 2 \cdot (a-1) \cdot e^{-a\alpha}$ we get the above expression. Similarly to the proof of Theorem 3.9, we get that

$$T(x) = -W(x) - \frac{1}{2}W^{2}(x),$$

with convergence within $|x| \leq e^{-1}$ [12].

Since the function $f(\alpha) = -\alpha \cdot 2 \cdot (a-1) \cdot e^{-a\alpha}$ gets its minimum at $\alpha = a^{-1}$, where it equals $-\frac{2(a-1)}{a}e^{-1}$, and $\left|-\frac{2(a-1)}{a}e^{-1}\right| \leq e^{-1}$ for all $a \in [1,2]$, then for all α we can substitute $x = -\alpha \cdot 2 \cdot (a-1) \cdot e^{-a\alpha}$. Hence, it is within the radius of convergence of T(x).

Finally, for the case where a = 1, then $d_2 = 0$ and $d_1 = n$. Therefore, the expression for the expected maximum matching size is reduced to $m - \left(m \cdot \left(1 - \frac{1}{m}\right)^n\right)$. Thus,

$$\gamma_a = \lim_{n \to \infty} \frac{\mu(G_a)}{n} = \lim_{n \to \infty} \frac{1}{n} \cdot \left(m - \left(m \cdot \left(1 - \frac{1}{m} \right)^n \right) \right) = \frac{1}{\alpha} - \frac{1}{\alpha} \cdot e^{-\alpha}.$$

A.10 Proof of Corollary 4.6

We show that γ_a is strictly monotonically increasing, thus $\gamma_a < 1$ for $1 \le a < 2$, since $\gamma_a = 1$ for a = 2. This is shown by differentiating γ_a with respect to a:

$$\frac{d\gamma_a}{da} = -\frac{1}{4\alpha^2 \left(a-1\right)^2} \cdot \left(W\left(-2\alpha \left(a-1\right) \cdot e^{-a\alpha}\right) + 2\alpha \left(a-1\right)\right) \cdot W\left(-2\alpha \left(a-1\right) \cdot e^{-a\alpha}\right)$$

Both the first factor $-\frac{1}{4\alpha^2(a-1)^2}$ and the third factor $W(-2\alpha (a-1) \cdot e^{-a\alpha})$ are negative. Thus, if the second factor is positive then $\frac{d\gamma_a}{da}$ is an increasing function with respect to $a \in [1, 2)$.

If $\alpha > 0.5$, then $2\alpha (a - 1) > 1$, and since W(x) is minimized for $x = -\frac{1}{e}$ where it equals -1, the second factor is positive. On the other hand, consider that $\alpha \le 0.5$. Since $W\left(-2\alpha (a - 1) \cdot e^{-2\alpha(a - 1)}\right) = -2(a - 1)\alpha$ and W(x) is an increasing function, then we have to show that $-2\alpha (a - 1) \cdot e^{-2\alpha(a - 1)} < -2\alpha (a - 1) \cdot e^{-a\alpha}$, that is, $-2\alpha (a - 1) > -a\alpha$. The last inequality can easily be shown for $1 \le a < 2$.

A.11 Proof of Theorem 4.7

We compute the limit of $\frac{\mu(G_p)}{n}$ as $n \to \infty$.

$$\gamma_p = \lim_{n \to \infty} \frac{\mu(G_p)}{n} = \lim_{n \to \infty} \frac{1}{n} \sum_{d_2 = 0}^n \binom{n}{d_2} \cdot p^{d_2} \cdot (1-p)^{n-d_2} \cdot \mu\left(G_{a=1+\frac{d_2}{n}}\right)$$

Let $X \sim Bin(n, p)$ be the random variable counting the number of vertices in L that choose 2 vertices in R. By summing over three disjoint ranges of possible values for d_2 , we get

$$\begin{split} \gamma_p &= \lim_{n \to \infty} \sum_{d_2=0}^{\lfloor np-n^{\frac{3}{4}} \rfloor} \Pr\left\{X = d_2\right\} \cdot \frac{1}{n} \cdot \mu\left(G_{a=1+\frac{d_2}{n}}\right) + \\ &\lim_{n \to \infty} \sum_{d_2=\lfloor np-n^{\frac{3}{4}} \rfloor+1}^{\lfloor np+n^{\frac{3}{4}} \rfloor-1} \Pr\left\{X = d_2\right\} \cdot \frac{1}{n} \cdot \mu\left(G_{a=1+\frac{d_2}{n}}\right) + \\ &\lim_{n \to \infty} \sum_{d_2=\lfloor np-n^{\frac{3}{4}} \rfloor}^{n} \Pr\left\{X = d_2\right\} \cdot \frac{1}{n} \cdot \mu\left(G_{a=1+\frac{d_2}{n}}\right) \end{split}$$

By Chebyshev's inequality we get that $\Pr\left\{|X - np| > n^{\frac{1}{4}}\sqrt{np(1-p)}\right\} \le \frac{1}{n^{\frac{1}{4}}}$. Since $p(1-p) \le 1$, we get that $\Pr\left\{|X - np| > n^{\frac{3}{4}}\right\} \le \frac{1}{n^{\frac{1}{4}}}$. By the fact that $\frac{1}{n} \cdot \mu\left(G_{a=1+\frac{d_2}{n}}\right) \le 1$, we find that the first and the third limits go to zero.

Since the function $\mu(G_a)$ is increasing with respect to a (this can be shown by a simple combinatorial argument), we get the following lower bound:

$$\lim_{n \to \infty} \left(1 - \frac{1}{n^{\frac{1}{4}}} \right) \cdot \frac{1}{n} \cdot \mu \left(G_{a=1 + \frac{\lfloor np - n^{\frac{3}{4}} \rfloor + 1}{n}} \right) \leq \lim_{n \to \infty} \sum_{d_2 = \lfloor np - n^{\frac{3}{4}} \rfloor + 1}^{\lfloor np + n^{\frac{3}{4}} \rfloor - 1} \Pr\left\{ X = d_2 \right\} \cdot \frac{1}{n} \cdot \mu \left(G_{a=1 + \frac{d_2}{n}} \right) = \gamma_p$$

as well as the following upper bound:

$$\gamma_p = \lim_{n \to \infty} \sum_{d_2 = \lfloor np - n^{\frac{3}{4}} \rfloor + 1}^{\lfloor np + n^{\frac{3}{4}} \rfloor - 1} \Pr\left\{ X = d_2 \right\} \cdot \frac{1}{n} \cdot \mu\left(G_{a=1 + \frac{d_2}{n}}\right) \le \lim_{n \to \infty} 1 \cdot \frac{1}{n} \cdot \mu\left(G_{a=1 + \frac{\lfloor np + n^{\frac{3}{4}} \rfloor - 1}{n}}\right).$$

By the squeeze theorem, we get the claimed result.

A.12 Proof of Theorem 5.3

Similarly to the proof of Theorem 3.6, our proof is based on counting the expected number of vertices in L that are not in some specific maximum matching M of G_{β} , based on the decomposition of G into its connected components. As in the proof of Theorem 3.6, we consider the number of connected components with exactly s vertices in L and q = s + 1 vertices in $R_u \cup R_d$, where we have to sum over all possible combinations (i, s + 1 - i), where i corresponds to the number of vertices taken from R_u and s + 1 - i corresponds to those taken from R_d .

Thus, the expected number of connected components in G_{β} with s vertices in L, i vertices in R_u and s + 1 - i vertices in R_d is given by:

$$\binom{n}{s} \cdot \binom{\beta \cdot m}{i} \binom{(1-\beta) \cdot m}{s+1-i} \cdot \left(1 - \frac{i}{\beta \cdot m}\right)^{n-s} \cdot \left(1 - \frac{s+1-i}{(1-\beta) \cdot m}\right)^{n-s} \cdot \left(\frac{i}{\beta \cdot m}\right)^s \cdot \left(\frac{s+1-i}{(1-\beta) \cdot m}\right)^s \cdot P_{i,s+1-i},$$

The above expression consists of the following factors (in order):

(*i*) choosing the s vertices in L;

(*ii*) choosing the *i* vertices in R_u ;

(*iii*) choosing the s + 1 - i vertices in R_d ;

(*iv*) the probability that all *i* vertices in R_u may be connected only to the chosen *s* vertices in *L*;

(v) the probability that all s + 1 - i vertices in R_d may be connected only to the chosen s vertices in L; (vi) the probability that all s vertices in L are only connected to the i vertices in R_u ;

(vii) the probability that all s vertices in L are only connected to the s + 1 - i vertices in R_d ; and, (viii) the probability that all chosen vertices are connected.

Finally, adding the expressions for all possible s's and i's and subtracting it from m yields the claimed result.

A.13 Proof of Theorem 5.4

As in the proof of Theorem 3.9, we compute the limit of $\frac{\mu(G)}{n}$ as $n \to \infty$. We consider the case where $\alpha = \frac{n}{m}$ and $0 \le \beta \le 1$ are fixed:

$$\gamma_{\beta} = \lim_{n \to \infty} \frac{\mu \left(G_{\beta}\right)}{n} = \lim_{n \to \infty} \frac{1}{n} \cdot \left(m - \sum_{s=0}^{n} \binom{n}{s} \cdot \sum_{i=b_{1}}^{b_{2}} \binom{\beta \cdot m}{i} \binom{(1-\beta) \cdot m}{s+1-i} \cdot \left(1 - \frac{i}{\beta \cdot m}\right)^{n-s} \cdot \left(\frac{i}{\beta \cdot m}\right)^{s} \cdot \left(\frac{s+1-i}{(1-\beta) \cdot m}\right)^{s} \cdot P_{i,s+1-i}\right)$$

By substituting the expression for $P_{i,s+1-i}$ from Theorem 5.3, and moving $\binom{n}{s}$ inside the second summation, we get:

$$\gamma_{\beta} = \lim_{n \to \infty} \left(\frac{1}{\alpha} - \frac{1}{n} \cdot \sum_{s=0}^{n} \sum_{i=0}^{s+1} \binom{n}{s} \cdot \binom{\beta \cdot m}{i} \binom{(1-\beta) \cdot m}{s+1-i} \cdot \left(1 - \frac{i}{\beta \cdot m}\right)^{n-s} \cdot \left(1 - \frac{s+1-i}{(1-\beta) \cdot m}\right)^{n-s} \left(\frac{i}{\beta \cdot m}\right)^{s} \cdot \left(\frac{s+1-i}{(1-\beta) \cdot m}\right)^{s} \cdot \frac{i^{(s+1-i)-1} \cdot (s+1-i)^{i-1} \cdot (i+(s+1-i)-1)!}{(i \cdot (s+1-i))^{i+(s+1-i)-1}}\right)$$

By substituting $\alpha = \frac{n}{m}$, we get:

$$\gamma_{\beta} = \lim_{n \to \infty} \left(\frac{1}{\alpha} - \frac{1}{n} \cdot \sum_{s=0}^{n} \sum_{i=0}^{s+1} \binom{n}{s} \cdot \binom{\frac{\beta}{\alpha} \cdot n}{i} \binom{\frac{1-\beta}{\alpha} \cdot n}{s+1-i} \cdot \left(1 - \frac{i}{\frac{\beta}{\alpha} \cdot n}\right)^{n-s} \cdot \left(1 - \frac{s+1-i}{\frac{1-\beta}{\alpha} \cdot n}\right)^{n-s} \left(\frac{i}{\frac{\beta}{\alpha} \cdot n}\right)^{s} \cdot \left(\frac{s+1-i}{\frac{1-\beta}{\alpha} \cdot n}\right)^{s} \cdot \frac{i^{(s+1-i)-1} \cdot (s+1-i)^{i-1} \cdot (i+(s+1-i)-1)!}{(i \cdot (s+1-i))^{i+(s+1-i)-1}}\right)$$

As in the proof of Theorems 3.9 and 4.5, using the monotone convergence theorem [39], we can put the limit inside the sum. By further simplifying the above expression with similar consideration to the proofs of Theorems 3.9 and 4.5, we get eventually:

$$\gamma_{\beta} = \frac{1}{\alpha} - \frac{\beta \cdot (1-\beta)}{\alpha^2} \sum_{s=0}^{\infty} \sum_{i=0}^{s+1} \frac{i^{(s+1-i)-1} \cdot (s+1-i)^{i-1}}{i! \cdot (s+1-i)!} \cdot \left(\frac{\alpha}{\beta} \cdot e^{-\frac{\alpha}{1-\beta}}\right)^{s+1-i} \cdot \left(\frac{\alpha}{1-\beta} \cdot e^{-\frac{\alpha}{\beta}}\right)^i$$

We switch the order of summation and get that $i \in \{0, 1, ...\}$ and s goes from $\max\{0, i-1\}$ to ∞ . We also substitute j = s + 1 - i (or s = i + j - 1). Thus,

$$\gamma_{\beta} = \frac{1}{\alpha} - \frac{\beta \cdot (1-\beta)}{\alpha^2} \sum_{i=0}^{\infty} \sum_{j=\max\{0,i-1\}}^{\infty} \frac{i^{j-1} \cdot j^{i-1}}{i! \cdot j!} \cdot \left(\frac{\alpha}{1-\beta} \cdot e^{-\frac{\alpha}{\beta}}\right)^i \cdot \left(\frac{\alpha}{\beta} \cdot e^{-\frac{\alpha}{1-\beta}}\right)^j \tag{4}$$

Let $T(x,y) = \sum_{j+i\geq 1} \frac{i^{j-1} \cdot j^{i-1}}{i! \cdot j!} \cdot x^i \cdot y^j$. This expression has been previously found [16] to be the multivariate formal power series about the point $(x_0, y_0) = (0, 0)$ of $t(x, y) = t_1(x, y) + t_2(x, y) - t_1(x, y) \cdot t_2(x, y)$ where $t_1(x, y)$ and $t_2(x, y)$ are given by the following implicit multivariate functions:

$$x = t_1(x, y) \cdot e^{-t_2(x, y)} \quad , \quad y = t_2(x, y) \cdot e^{-t_1(x, y)}$$
(5)

However, the mentioned range of convergence in [16] is insufficient for our case. (Note also that in [16] the sums should be over $i + j \ge 1$ and not over $i, j \ge 0$.)

Since we compute the limit normalized expected maximum matching, then the expression for γ_{β} in Equation (4) is bounded from below by 0, thus, by Equation (4) the double summation is bounded from above by a constant. On the other hand, all terms in the summation in Equation (4) are positive. Then, if we look at the partial-sum series (by defining an arbitrary order), we get an increasing series which is bounded. Thus, by the monotone convergence theorem the double series converges for any values x and y satisfying $x = \frac{\alpha}{1-\beta} \cdot e^{-\frac{\alpha}{\beta}}$ and $y = \frac{\alpha}{\beta} \cdot e^{-\frac{\alpha}{1-\beta}}$.

However, the multivariate functions in Equation (5) have multiple branches (as the Lambert-W function does [12]), that is, for a given x and y there is more than one solution. We aim to find this branch in terms of t_1 and t_2 . We use the implicit function theorem to find the derivatives singularities. The Jacobian is given by

$$J = \begin{pmatrix} e^{-t_2(x,y)} & -t_1(x,y) \cdot e^{-t_2(x,y)} \\ -t_2(x,y) \cdot e^{-t_1(x,y)} & e^{-t_1(x,y)} \end{pmatrix},$$

and it is invertible wherever $|J| \neq 0$. Thus, there is a derivative singularity in case $t_1(x, y) \cdot t_2(x, y) = 1$, which is the only solution. Therefore, as the given formal power series in Equation (4) is about the point $(x_0, y_0) = (0, 0)$ (which corresponds to $\alpha = 0$), where $t_1 = t_2 = 0$, it converges to the branch where $t_1(x, y) \cdot t_2(x, y) \leq 1$ (note that both $t_1(x, y)$ and $t_2(x, y)$ are always positive).

A.14 Proof of Theorem 6.1

We first establish a few lemmas before proving the result. As before, we start by considering a deterministic bipartite graph $H = \langle L_H + R_H, E_H \rangle$ with degree d of each vertex in L_H , where $|L_H| = s$ and $|R_H| = q$.

Lemma A.1. If $(d-1) \cdot s \leq q-2$, then H is not connected.

Proof. As in the proof of Lemma 3.1, the proof follows by induction on s. For s = 1, there are d edges in the graph and therefore every graph with $q \ge d + 1$ is not connected. Assuming that the claim holds up until s = s', we next prove that it holds for any bipartite graph H' such that $|L_{H'}| = s' + 1$ and $|R_{H'}| \ge (d-1) \cdot (s'+1) + 2$. Assume towards a contradiction that there is a graph H' which is connected.

We first show that there are d-1 vertices $v_{r_1}, v_{r_2}, \ldots, v_{r_{d-1}}$ in $R_{H'}$, all of a degree 1 such that they are connected to the same vertex $v_{\ell} \in R_{H'}$: The sum of right-side vertex degree is $d \cdot (s'+1)$. Also, since the graph is connected there are no right-side vertices with degree 0. This implies that there are at least $(d-2) \cdot (s'+1) + 2$ vertices of degree 1, thus there exists a vertex $v_{\ell} \in R_{H'}$ as claimed.

By the induction hypothesis, the graph induced by $L_{H'} \setminus \{v_\ell\}$ and $R_{H'} \setminus \{v_{r_1}, v_{r_2}, \ldots, v_{r_{d-1}}\}$ is not connected, which implies that it has at least two connected components. In H', v_ℓ is connected to all vertices $v_{r_1}, v_{r_2}, \ldots, v_{r_{d-1}}$. Since its degree is d it can be connected only to one of these components. This implies that H' is not connected as well, and the claim follows.

Lemma A.2. If H is connected and $(d-1) \cdot s = q-1$ then $\mu(H) = s$.

Proof. Assume towards a contradiction that $\mu(H) < s$, and consider some maximum matching M. Let $v_{\ell} \in L_H$ be a vertex that is not in the maximum matching M, and $v_{r_1}, v_{r_2}, \ldots, v_{r_{d-1}}$ be the vertices in R (which are not necessarily distinct) that are connected to v_{ℓ} . All vertices $v_{r_1}, v_{r_2}, \ldots, v_{r_{d-1}}$ are connected also to another vertex in L_H , otherwise v_{ℓ} was in the maximum matching M.

Consider the bipartite graph $\hat{H} = \langle \hat{L}_H + \hat{R}_H, \hat{E}_H \rangle$, which is given by removing v_ℓ from H. Since the right-side vertices $v_{r_1}, v_{r_2}, \ldots, v_{r_{d-1}}$ are also connected to the other left-side vertices (except v_ℓ), the bipartite graph \hat{H} is connected. However, we get that $|\hat{L}_H| = s - 1$ and $|\hat{R}_H| = (d - 1) \cdot s + 1$, which contradicts with Lemma A.1.

We note that in contrast to Lemma 3.2, the corresponding proposition is not true for d > 2; that is, if H is connected and $s \le q$, then the maximum matching size is not necessarily s. As a counter example, consider the case where d = 3 and s = q = 3, where two left-side vertices choose the same single right-side vertex (using all their 3 choices), and the other left-side vertex chooses all 3 right-side vertices. The resulting bipartite graph is clearly connected, but the maximum matching size is only 2 (only one of the first two left-vertices can be in the matching).

Lemma A.3. If $(d-1) \cdot s = q-1$ then H is connected if and only if it is a tree.

Proof. The proof consists of the exact same construction \hat{H} as in the proof of Lemma 3.4, where we eventually get a contradiction with Lemma A.1.

Lemma A.4. The number T_s^d of connected bipartite graphs H whose $|L_H| = s$ and $|R_H| = 2 (d-1) \cdot s + 1$ is $T_s^d = \frac{((d-1)\cdot s+1)!}{((d-1)!)^s} ((d-1)\cdot s+1)^{s-2}$.

Proof. By Lemma A.3, we have to count the number of bipartite trees over the two disjoint sets L_H and R_H of size s and $(d-1)\cdots + 1$. Since H is a tree, then there are no cycles. Consequently, each one of the vertices in L_H is connected to d distinct vertices in R_H . Moreover, no two vertices in L_H share



Figure 1: the Lambert-W function

more than 1 vertex in R_H . For each vertex $v_{\ell} \in L_H$, let S_v be the set of the d right-side vertices that v_{ℓ} is connected to and also let the cycle $C_{v_{\ell}}$ be a cycle that consists of the d vertices of S_v .

Consider the graph $\hat{H} = \langle \hat{R}_H, \hat{E}_H \rangle$, which is given by connecting each cycle $C_{v_{\ell_1}}$ to $C_{v_{\ell_2}}$ using a common vertex v_r if and only if v_r is connected to both v_{ℓ_1} and v_{ℓ_2} . The resulting graph \hat{H} is a Husimi graph over $(d-1)\cdot s+1$ vertices, where the number of such (labeled) graphs is $\frac{((d-1)\cdot s+1)!}{((d-1)!)^{s}\cdot s!} ((d-1)\cdot s+1)^{s-2} [7,22].$ Finally, each set S_v is determined by the (labeled) vertex in R_L . Thus, we multiply by s! the above

expression.

We are now able to prove the result.

Proof. [of Theorem 6.1] Let M be a maximum matching of G. Similarly to the proof of Theorem 3.6, the proof is based on counting the expected number of vertices in R that are not part of M, and on the decomposition of G into its connected components.

We count the expected number of connected components with s left-side vertices and q = (d-1). s+1 right-side vertices. By Lemma A.2, the maximum matching size of each such connected component is exactly s. Thus, there are q - s right-side vertices that are not in M.

Let H be a bipartite graph $H = \langle L_H + R_H, E_H \rangle$, with degree d for all vertices in L_H , where $|L_H| = s$ and $|R_H| = q$. The probability P_s that H is connected is given by $P_s = \frac{(d!)^s T_s^d}{q^{d \cdot s}}$.

The remainder of the proof is similar to the proof of Theorem 3.6.

B **Evaluation**

B.1 The Lambert-*W* **Function**

The Lambert-W function, usually denoted by $W(\cdot)$, is given by the following implicit representation:

$$z = W(z) \cdot e^{W(z)}$$

where z is a complex number [12].

For real valued arguments, i.e. z is real valued, W(z) has two real-valued branches: the principal branch, denoted by $W_0(\cdot)$ and the branch $W_{-1}(\cdot)$. Figure 1 shows the two real-valued branches. For instance, $W_0(-e^{-1}) = W_{-e^{-1}} = -1$ and $W_0(0) = 0$.



Figure 2: Expected maximum matching size for various values of n and m (normalized by n)



Figure 3: Limit expected maximum matching size for various values of load α , normalized by n

B.2 Expected Maximum Matching Size With d = 2

Figure 2 shows the expected maximum matching size normalized by n for various values of n and m. We show the expected maximum matching size both via our analytical model from Theorem 3.6 and via simulations. For each instance of n and m, we randomized 10000 bipartite graphs. The results fairly confirm that our model is accurate, and also show the convergence of the expected maximum matching size to its limit. A simple evaluation appears in the following example.

Example B.1. In case n = m = 2 (and d = 2), the expected maximum matching size is $\mu(G) = \frac{15}{8} = 1.875$. This simple result can be justified as follows: In all cases the maximum matching size is 2, except for the two cases of maximum matching of size 1, where all 4 edges are connected to a specific vertex in R. Each such case occurs with probability $\left(\frac{1}{2}\right)^4$. Hence, $\mu(G) = 2 - \frac{1}{16} - \frac{1}{16} = \frac{15}{8}$.

Figure 3 shows the expected maximum matching size normalized by n as found in Theorem 3.9, for various values of load α , both via our analytical model and via simulations. The simulations were performed using m = 1000 and $n = \alpha \cdot m$. For each value of α , we randomized 100 bipartite graphs. The results fairly confirm that our model is accurate.

We conclude by the following simple example:

Example B.2. In case $\alpha = 1$, that is n = m, the normalized limit expected maximum matching size is

$$\gamma = 1 + \frac{1}{2} \cdot W\left(-2 \cdot e^{-2}\right) + \frac{1}{4}W^2\left(-2 \cdot e^{-2}\right) \approx 0.8381.$$



Figure 4: Limit expected maximum matching size for various values of α and a, normalized by n



Figure 5: Limit expected maximum matching size for various values of β and α , normalized by n

B.3 Expected Maximum Matching Size With $d_v \leq 2$

Figure 4 shows the normalized limit expected maximum matching size, for various values of load α and average number of choices a, both via our analytical model (from Theorem 3.9) as well as via simulations. The simulations were performed using m = 1000 and $n = \alpha \cdot m$, where for each instance of the simulation we randomized 100 bipartite graphs. The results fairly confirm that our model is accurate.

B.4 Expected Maximum Matching Size With Static Partition

Figure 5 shows the limit expected maximum matching size normalized by n, for various values of load α and partition β , both via our analytical model (from Theorem 5.4) and via simulations. The simulations were performed using m = 1000 and $n = \alpha \cdot m$. For each pair of values of α and β , we randomized 100 bipartite graphs. The results fairly confirm that our model is accurate.

As expected, the limit expected maximum matching size is symmetric around $\beta = 0.5$. In case $\alpha = 0.5$ and $\beta < 0.5$, while it seems that the normalized limit expected maximum matching size is 1, it is not the case. For instance, in case $\alpha = 0.5$ and $\beta = 0.45$, we get that $1 - \gamma_{\beta} \approx 1.675 \cdot 10^{-7}$. However, there are cases where imbalance in the partition sizes does not reduce γ_{β} , as shown for instance in Corollary 5.6.



Figure 6: Upper bound on the normalized expected maximum matching size for $\alpha = 1$ as a function of d

B.5 Expected Maximum Matching Size With d > 2

We evaluate the upper bound found for the expected matching size (Theorem 6.1). Figure 6 shows our upper bound as well as simulation results for various values of the number of choices d. We took n = m = 100, while for each instance of d, we randomized 10^5 bipartite graphs. In the case of d = 2, our upper bound matches the exact expression found in Theorem 3.6 and thus matches the simulation results. In addition, we can compare simulation results for higher values of d with our bounds. For instance, in the case of d = 3 the normalized expected maximum matching size via the simulation is 0.9402, while our upper bound is 0.9508. In case d = 4, we get a simulation value of 0.9795, while the corresponding upper bound is 0.9820.