Improved Algorithm for Dynamic b-Matching

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Abstract

Recently there has been extensive work on maintaining (approximate) maximum matchings in dynamic graphs. We consider a generalisation of this problem known as the *maximum b-matching*: Every node v has a positive integral capacity b_v, and the goal is to maintain an (approximate) maximum-cardinality subset of edges that contains at most b_v edges incident on every node v. The maximum matching problem is a special case of this problem where $b_v = 1$ for every node v.

Bhattacharya, Henzinger and Italiano [ICALP 2015] showed how to maintain a $O(1)$ approximate maximum b-matching in a graph in $O(\log^3 n)$ amortised update time. Their approximation ratio was a large (double digit) constant. We significantly improve their result both in terms of approximation ratio as well as update time. Specifically, we design a randomised dynamic algorithm that maintains a $(2+\epsilon)$-approximate maximum b-matching in expected amortised $O(1/\epsilon^4)$ update time. Thus, for every constant $\epsilon \in (0,1)$, we get expected amortised $O(1)$ update time. Our algorithm generalises the framework of Baswana, Gupta, Sen [FOCS 2011] and Solomon [FOCS 2016] for maintaining a maximal matching in a dynamic graph.

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1 Introduction

In dynamic graph algorithms, we want to maintain a solution to an optimisation problem on an input graph that is undergoing a sequence of edge insertions/deletions. The time taken to modify the solution after an *update* (edge insertion/deletion) is called the *update time* of the dynamic data structure. The challenge is to design dynamic data structures whose update times are significantly faster than recomputing the solution from scratch after each update using the best static algorithm. In recent years, there has been extensive work on dynamic graph algorithms for (approximate) maximum matching [10, 1, 12, 8, 3, 9, 2, 11, 6, 5]. A matching $M \subseteq E$ in a graph $G = (V, E)$ is a subset of edges that do not share a common endpoint. In this problem, the goal is to maintain a matching of (approximately) maximum size in an input graph undergoing a sequence of edge insertions/deletions.

The first significant result on dynamic matching was due to Onak and Rubinfeld [10], who gave a randomised data structure with approximation ratio $O(1)$ and amortised update time of $O(\log^2 n)$. Baswana, Gupta and Sen [1] improved this approximation ratio to 2 and the amortised update time to $O(\log n)$. Very recently, in a breakthrough result Solomon [12] built upon the algorithmic framework of Baswana, Gupta and Sen [1] to present a randomised data structure with approximation ratio 2 and *constant amortised update time*. This is the


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first paper to achieve constant (and hence optimal) update time for *any* graph problem. All the data structures described so far are randomised. There are deterministic data structures for this problem, with $(2+\epsilon)$-approximation ratio and $O(\text{poly}\log n)$ amortised update time, that are due to Bhattacharya, Henzinger and Italiano [5] and Bhattacharya, Henzinger and Nanongkai [6]. They show how to maintain a $(2+\epsilon)$-approximate maximum *fractional* matching in $O(\log n/\epsilon^2)$ amortised update time. Then they *round* these fractional weights deterministically in a dynamic setting to get an integral matching.

We focus on a generalisation of the dynamic matching problem. We are given an input graph $G=(V,E)$ with $n=|V|$ nodes. At each time-step, an edge is inserted into/deleted from the graph. Each node $v \in V$ has a positive integral *capacity* b_v. The node-set V and the capacities $\{b_v\}$ remain unchanged over time. A b-matching is a subset of edges $\mathcal{M} \subseteq E$ that contains at most b_v edges incident on every node $v \in V$. Note that if $b_v = 1$ for every node $v \in V$, then $\mathcal{M}$ is a matching. Our goal is to maintain a b-matching of (approximately) maximum size in this dynamic graph G. This problem was first considered by Bhattacharya, Henzinger and Italiano [4]. Extending the dynamic data structure for fractional matching [5], they first showed how to maintain a fractional b-matching. Then they randomly sampled the edges from this fractional b-matching, with probabilities proportional to their edge-weights, and got an integral b-matching. This leads to a randomised data structure with a (large, double digit) constant approximation ratio and $O(\log^3 n)$ amortised update time with high probability. Recently, Gupta et. al [7] improved the update time in [4] for the set cover problem. However, their result does not imply our b-matching result.

Our result. We significantly improve upon the previous result on dynamic b-matching [4], both in terms of the approximation ratio and update time. Specifically, we present a randomised data structure with approximation ratio $(2+\epsilon)$ and expected amortised update time $O(1/\epsilon^4)$. Note that our update time does *not* depend on the capacities $\{b_v\}$. On the other hand, similar to the previous paper [4], we assume an oblivious adversary.

Our technique. We build upon the work of Baswana, Gupta, Sen [1] and Solomon [12]. Before proceeding any further, we briefly review their framework for dynamic matching. A matching $M \subseteq E$ is *maximal* if every unmatched edge $(u,v) \in E \setminus M$ has at least one matched endpoint. The size of a maximal matching is a 2-approximation to the size of the maximum matching. Note that we can easily maintain a maximal matching in a dynamic graph in $O(d)$ update time, where d is an upper bound on the maximum degree of a node. This holds since after the insertion/deletion of an edge (u,v), we only need to check the neighbours of u and v to preserve maximality, and there are at most $O(d)$ such neighbours. To improve the update time, the papers [1, 10, 12] use the following idea. We pick a neighbour y of x uniformly at random from the set N_x, where N_x is the set of all neighbours of x, and add the edge (x,y) to the matching M. If the sequence of edge insertions/deletions in G is oblivious to the random bits used by the algorithm, then in expectation this edge (x,y) will not be deleted from G before half of the current edges in N_x gets deleted. And when such an edge $(x,z) \in N_x \setminus \{(x,y)\}$ gets deleted, the algorithm does not need to perform any computation for the node x (since the node remains matched in M). Thus, although the algorithm takes $O(|N_x|)$ time to find a new *mate* for x, it does not need to do anything for the next $|N_x|/2$ edge deletions incident on x, in expectation. This helps bring down the amortised update time to $O(1)$. This is the high level idea behind the dynamic algorithms in [1, 12]. The actual algorithms, however, are significantly more complicated, for the following reason. In the preceding summary, we said that the node x picks a neighbour y uniformly at random from

the set N_x. But what if the node y itself is matched to some other node z (i.e., $(y,z) \in M$)? In that case, in order to add the edge (x,y) to the matching M, we first need to unmatch the edge (y,z) by removing it from M. But this means that we will now need to find a new mate for z. In general, this can lead to a long chain of edges alternately being matched and unmatched. To address this concern, the papers [1, 12] construct a *hierarchical partition* of the node-set into $O(\log n)$ levels. Suppose that a node x is at level i, and want to find a new mate for x. The papers [1, 12] ensure that x has roughly α^i many neighbours at strictly lower levels $\{0, \ldots, i-1\}$, for some constant $\alpha > 1$. We now pick a node y uniformly at random *only from* these neighbours of x at levels $j < i$. The papers further ensure that every matched edge has both its endpoints at the same level. Hence, the node (say z) that is matched to y will have $\ell(z) = \ell(y) < i = \ell(x)$. We now match the edge (x,y), bring the node y up to level i, and unmatch the edge (y,z). The node z now has to find a new mate for itself. But note that the level of z is strictly less than the level of x. Hence, this chain of alternate matchings and unmatchings of edges cannot go on for more than $O(\log n)$ levels.

In order to extend the above framework to maximum b-matching, we have to overcome several technical difficulties. First, since a node can have multiple matched edges incident on it in a b-matching, we can no longer ensure that both the endpoints of every matched edge are on the same level. Instead, we maintain an invariant which states that if a node v has b_v matched edges incident on it, then at least one of these matched edges, say (u,v), must have its other endpoint u at a level that is not larger than the level of v (see Invariant 3). Second, unlike the papers [1, 12], we can no longer ensure that if a node z becomes unmatched while we are finding a new mate for a different node x, then $\ell(x) > \ell(z)$. Indeed, there are instances in our algorithm where both x and z can be at the same level, and we need to show that this situation does not occur too often. Finally, in the papers [1, 12], at any point in time there can be multiple nodes in the hierarchical partition that are *dirty*, meaning that they violate one of the invariants. The algorithms in [1, 12] run a WHILE loop, and each iteration of the WHILE loop picks *any arbitrary* dirty node and restores all the invariants it ought to satisfy (this can lead to other nodes becoming dirty). In sharp contrast, the WHILE loop in our dynamic algorithm has to pick the dirty nodes *in a specific order*, preferring one type of dirty nodes over another. This preferential ordering among the dirty nodes turns out to be crucial in analysing the amortised update time of our algorithm. See Sections 2 and 3 for details.

2 Our Dynamic Algorithm for Maximum b-Matching

Let $G = (V, E)$ denote the input graph whose edge-set E changes dynamically. Every node $v \in V$ has a positive integral *capacity* b_v associated with it. The node-set V and the capacities $\{b_v\}$ do not change over time. Let $n = |V|$ be the number of nodes in G. A subset of edges $\mathcal{M} \subseteq E$ is a *b-matching* iff for all nodes $v \in V$, we have $|\{(u,v) \in \mathcal{M} \mid (u,v) \in E\}| \leq b_v$. We say that an edge $e \in E$ is *matched* in $\mathcal{M}$ iff $e \in \mathcal{M}$.

Throughout this paper, we fix any constant $\epsilon \in (0, 1/2)$ and define $\alpha = 5/\epsilon$. We will maintain a partition of the node-set V into $L+2$ *levels* $\{-1, 0, \ldots, L\}$, where $L = \lceil \log_\alpha n \rceil$. Let $\ell(v) \in \{-1, \ldots, L\}$ denote the level of a node v. The value of $\ell(v)$ changes over time. In our dynamic algorithm, whenever a node v moves to a level j, we will require that it has at most $2b_v \cdot \alpha^{j+1}$ neighbours u with $\ell(u) \leq j$. Further, any node v at level -1 has at most $O(b_v \cdot \alpha^0) = O(b_v)$ many neighbours at level -1.

The *level of an edge* is the maximum level among its endpoints, i.e., $\ell(u,v) = \max(\ell(u), \ell(v))$ for all $(u,v) \in E$. For every node $v \in V$ and every level $j \in [-1, L]$, we let $\mathcal{E}_v^j = \{(u,v) \in E : \ell(u,v) = j\}$ be the set of edges incident on v that are at level j. Since $\ell(u,v) \geq \ell(v)$ for all

edges $(u, v) \in E$, it follows that $\mathcal{E}_v^j = \emptyset$ for all levels $j < \ell(v)$. We say that an edge $(u, v) \in E$ is *owned by* its endpoint v iff $\ell(u) < \ell(v)$. Equivalently, we say that v is the *owner* of the edge (u, v). We let $\mathcal{O}_v = \{(u, v) \in E : \ell(u) < \ell(v)\}$ denote the set of edges owned by $v \in V$ (Note that if both the end-points are at the same level then no vertex owns the edge). In a b-matching $\mathcal{M} \subseteq E$, we let $\mathcal{M}_v = \{(u, v) \in \mathcal{M}\}$ denote the set of all matched edges incident on a node $v \in V$. For any level $i \in [-1, L]$, we define $\mathcal{M}_v(i) = \{(u, v) \in \mathcal{M}_v : \ell(u, v) = i\}$ as the set of all matched edges incident on a node v at level i. We say that a node $v \in V$ is FULL in a b-matching $\mathcal{M} \subseteq E$ iff $|\mathcal{M}_v| = b_v$, and v is DEFICIENT in $\mathcal{M}$ iff $|\mathcal{M}_v| < (1 - \epsilon)b_v$. The node v is NON-DEFICIENT in $\mathcal{M}$ iff $|\mathcal{M}_v| \geq (1 - \epsilon)b_v$. Finally, for every node $v \in V$, if v is FULL, then we define $\text{BASE}(v)$ to be the smallest level j for which there is a matched edge $(u, v) \in \mathcal{M}_v$ at level $\ell(u, v) = j$. Else if v is not FULL, then we set $\text{BASE}(v) = \ell(v)$.

We maintain a b-matching $\mathcal{M} \subseteq E$ in G satisfying the three invariants below. The approximation guarantee of the algorithm follows from the first two invariants (see Theorem 4). Invariant 3 will be useful in bounding the amortised update time. The main result in this paper follows from Theorems 4, 5.

▶ **Invariant 1.** *Every node $v \in V$ at level $\ell(v) \geq 0$ is* NON-DEFICIENT.

▶ **Invariant 2.** *Every unmatched edge with level -1 has at least one* NON-DEFICIENT *endpoint.*

▶ **Invariant 3.** *For every node $v \in V$, we have $|\{(u, v) \in \mathcal{M} : \ell(u) > \ell(v)\}| < b_v$.*

We will assume that the initial graph is empty. So, all the three invariants hold. We now compare our invariants with those in ([1], [12]), that is in the maximal matching case when $b_v = 1$ for all $v \in V$. NON-DEFICIENT vertices are *free* vertices in this simple case. The first invariant then say that all the vertices at level ≥ 0 are matched . The second invariant says that each un-matched edge at level -1 has both endpoints free. In ([1],[12]), there can be no edge at level -1, since only *free* vertices settle at level -1. That is why the corresponding invariant for maximal matching is that *there is no edge with level* -1. We have generalised this invariant for b-matching. The last invariant implies that if at any point of time all the other endpoints of the matched edges incident on v have levels greater than v, then v has to move to a higher level. In the simple maximal matching case, this implies that end-points of the matched edges should always be at the same level. The reader can check that these three invariants are maintained in ([1],[12]), and here we generalise them for b-matching.

▶ **Theorem 4.** *Invariants 1, 2 imply that $\mathcal{M}$ is a $(2 + \epsilon)$-approximate maximum b-matching.*

Proof. (Sketch) Invariants 1, 2, ensure that $\mathcal{M}$ is *almost maximal*, as every unmatched edge has one endpoint x with $|\mathcal{M}_x| \geq (1 - \epsilon)b_x$. This implies $(2 + \epsilon)$-approximation. ◀

▶ **Theorem 5.** *There is a randomised algorithm that maintains a b-matching in a dynamic graph satisfying Invariants 1, 2, 3. It has an amortised update time of $O(1/\epsilon^4)$ in expectation.*

2.1 Handling the insertion/deletion of an edge

Data structures. Each node $x \in V$ maintains the edge-sets $\mathcal{M}_x, \mathcal{M}_x(j), \mathcal{E}_x^j$ and $\mathcal{O}_x$ as doubly linked lists, and the node also maintains the sizes of these sets. From the size of $\mathcal{M}_x$, we can infer whether the node x is DEFICIENT, NON-DEFICIENT or FULL. With appropriate pointers, an edge can be inserted into or deleted from a linked list in $O(1)$ time. Further, each node $x \in V$ maintains its level $\ell(x)$ and the value of $\text{BASE}(x)$.

```
01. WHILE the set of DIRTY nodes is nonempty
02.     IF the set of FULL-FROM-ABOVE nodes is nonempty, THEN
03.         Let x be a FULL-FROM-ABOVE node with the largest value of BASE(x).
04.         Call the subroutine FIX-DIRTY(x).   // See Section 2.1.1.
05.     ELSE
06.         Let x be any DIRTY node that is DEFICIENT at level ℓ(x) ≥ 0.
07.         Call the subroutine FIX-DIRTY(x).   // See Section 2.1.2.
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Figure 1 Fixing the dirty nodes.

Insertion/deletion of an edge (u, v). When an edge (u, v) is inserted into or deleted from the graph $G = (V, E)$, we first update the relevant data structures in $O(1)$ time. For the time being, to highlight the main idea behind our algorithm, we assume that the insertion/deletion of the edge (u, v) does not lead to a violation of Invariant 2 (which pertains to the subgraph induced by the nodes at level -1). In Section 2.1.4, we will show how to handle the nodes in level -1 and how to maintain Invariant 2. Right now we focus on maintaining the remaining two invariants. Accordingly, suppose that after the insertion/deletion of the edge (u, v), at least one of its endpoints violates Invariant 1 or 3. In general, if a node $x \in V$ violates Invariant 1 or 3, then we say that the node x is DIRTY. A dirty node is either DEFICIENT (if it violates Invariant 1) or FULL-FROM-ABOVE (if it violates Invariant 3). Thus, a node x is FULL-FROM-ABOVE iff $\text{BASE}(x) > \ell(x)$ and $|\mathcal{M}_x| = b_x$. Due to the insertion/deletion of an edge, its endpoints can become DIRTY. To ensure that no node remains DIRTY, we call the following subroutine in Figure 1.

We highlight two important aspects of this subroutine. First, during an iteration of the WHILE loop in Figure 1, the node x can move to a new level due to the call to FIX-DIRTY(x). Immediately after the call to FIX-DIRTY(x), the node x is no longer DIRTY. But the call to FIX-DIRTY(x) might lead to some other nodes becoming DIRTY, and these newly DIRTY nodes will be handled in subsequent iterations of the WHILE loop. Second, the WHILE loop in Figure 1 always gives preference to the FULL-FROM-ABOVE nodes over the DIRTY nodes that are DEFICIENT at a nonnegative level. Furthermore, among the FULL-FROM-ABOVE nodes, the WHILE loop picks the one with the largest BASE(.) value. The WHILE loop picks a DIRTY and DEFICIENT node only if there is no FULL-FROM-ABOVE node. This aspect of our algorithm will ensure that Lemma 9 holds, which, in turn, will play a crucial role in the analysis of the amortised update time.

2.1.1 The subroutine FIX-DIRTY(x) on a Full-from-above node x

Suppose that the subroutine FIX-DIRTY(x) is called on a FULL-FROM-ABOVE node x at level $\ell(x) = i$ with $\text{BASE}(x) = j$ (say). Since the node x is FULL-FROM-ABOVE, we have $j > i$. We first move the node x up from level i to level j, and update all the relevant data structures (as described in the beginning of Section 2.1). We next check the number of edges in $\mathcal{E}_x^j$, and, accordingly, we consider two cases.

Case 1. $|\mathcal{E}_x^j| \leq 2b_x \cdot \alpha^{j+1}$. In this case, the node x stays at level j and we terminate the subroutine. At this stage the node x has $|\mathcal{M}_x| = b_x$ and $\text{BASE}(x) = j = \ell(x)$. Hence, the node x is no longer DIRTY.

Case 2. $|\mathcal{E}_x^j| > 2b_x \cdot \alpha^{j+1}$. In this case, we find the *minimum* level $k \in [j+1, L]$ such that the number of edges (x, y) with $\ell(y) \leq k$ lies in the range $(2b_x \cdot \alpha^k, 2b_x \cdot \alpha^{k+1}]$. Such a level exists since $b_x \cdot \alpha^{L+1} \geq n >$ degree of x. We now move the node x from level j to level k. In doing so, x becomes the owner of all the edges whose other endpoints are at level $< k$ and we update all the relevant data structures. Next, we *unmatch* all edges $(x, y) \in \mathcal{M}_x$ whose other endpoints are at levels $\ell(y) < k$, and update all the relevant data structures. Let λ_x be the value of $|\mathcal{M}_x|$ at this stage. We now select $(b_x - \lambda_x)$ edges from $\mathcal{O}_x$ uniformly at random, and add them to $\mathcal{M}$ by calling RANDOM-SETTLE(x)(See Section 2.1.3). At this stage we have $|\mathcal{M}_x| = b_x$ and BASE(x) $= k = \ell(x)$, and hence x is no longer DIRTY. We terminate the subroutine at this point. This procedure can lead to other nodes becoming DIRTY, for two reasons. (1) When we unmatch an edge (x, y) with $\ell(y) < k$, the node y might become DEFICIENT and DIRTY. (2) Suppose that while executing RANDOM-SETTLE(x), we add an edge $(x, y) \in \mathcal{O}_x$ to $\mathcal{M}$. Then the node y can become FULL-FROM-ABOVE (and DIRTY). Else, it might happen that $|\mathcal{M}_y|$ becomes equal to $(b_y + 1)$ after we add the edge (x, y) to $\mathcal{M}$. Then we will need to unmatch some edge $(y, z) \in \mathcal{M}_y$, so as to ensure that $\mathcal{M}$ remains a valid b-matching. But this might lead to z becoming DEFICIENT and DIRTY. These newly DIRTY nodes will be handled in subsequent iterations of the WHILE loop in Figure 1.

▶ **Lemma 6.** *Suppose that we call FIX-DIRTY(x) on a* FULL-FROM-ABOVE *node x, and this moves the node x up from level i to level $k > i$. Then it takes $O(b_x \cdot \alpha^{k+1})$ time.*

Proof (Sketch). Only the edges $(x, y) \in E$ with $\ell(y) \leq k$ get affected as the node x moves to level k. The data structure associated with every other edge remains unchanged. Thus, the total runtime of the subroutine is proportional to the number of edges in $\{(x, y) \in E : \ell(y) \leq k\}$, plus the time taken by the potential call to RANDOM-SETTLE(x). The latter quantity is bounded in Lemma 10. It is easy to check that the former quantity is at most $2b_x\alpha^{k+1}$, for otherwise the node x would have moved up to a level larger than k. ◀

2.1.2 The subroutine FIX-DIRTY(x) on a Deficient node x

Suppose that the subroutine FIX-DIRTY(x) is called on a DEFICIENT node x at level $\ell(x) = j \geq 0$. We first check the number of edges in $\mathcal{E}_x^j$. Accordingly, we consider two cases.

Case 1. $|\mathcal{E}_x^j| > 2b_x \cdot \alpha^{j+1}$. Here, we perform the same steps as in Case 2 in Section 2.1.1.

Case 2. $|\mathcal{E}_x^j| \leq 2b_x \cdot \alpha^{j+1}$. In this case, we try to find a maximal level k in range $[0, j]$ such that the number of edges (x, y) with $\ell(y) < k$ lies in the range $(2b_x \cdot \alpha^k, 2b_x \cdot \alpha^{k+1}]$. If no such level exists, then we set $k = -1$. Next, we move the node x from level j to level k. In doing so, x dis-owns all the edges (and the other endpoints becomes the owner) that are at level $[k, j]$ and we update the relevant data structures. We then unmatch all the edges (x, y) whose other endpoints are at levels $\ell(y) < j$, and update the relevant data structures. Let λ_x be the value of $|\mathcal{M}_x|$ at this stage. If $k \geq 0$, then we call RANDOM-SETTLE(x), which selects $(b_x - \lambda_x)$ edges uniformly at random from $\mathcal{O}_x = \{(x, y) \in E : \ell(y) < k\}$, and adds those edges to $\mathcal{M}$. At this point $|\mathcal{M}_x| = b_x$ and BASE(x) $= k = \ell(x)$, and hence x is no longer DIRTY. So we terminate the subroutine. If $k = -1$, then instead of calling RANDOM-SETTLE(x), we follow the procedure in Section 2.1.4.

▶ **Lemma 7.** *Suppose that we call FIX-DIRTY(x) on a* DEFICIENT *node x at level $j \geq 0$. If the subroutine moves the node x to a level $k > j$, then it takes $O(b_x \cdot \alpha^{k+1})$ time.*

Proof. Exactly similar to the proof of Lemma 6. ◀

01. WHILE $|\mathcal{M}_x| < b_x$
02. Pick a uniformly random edge $(x, y) \in \mathcal{O}_x \setminus \mathcal{M}_x$, and add it to the b-matching. Specifically, set $\mathcal{M} \leftarrow \mathcal{M} \cup \{(x, y)\}$, and update the relevant data structures.
03. IF $|\mathcal{M}_y| = b_y + 1$, THEN
04. Select an edge $(y, z) \in \mathcal{M}_y \setminus \{(x, y)\}$ which minimises the value of $\ell(y, z)$, and unmatch the edge (y, z). Specifically, set $\mathcal{M} \leftarrow \mathcal{M} \setminus \{(y, z)\}$, and update the relevant data structures.

Figure 2 RANDOM-SETTLE(x).

▶ **Lemma 8.** *Suppose that we call FIX-DIRTY(x) on a* DEFICIENT *node x at level $j \geq 0$. If the subroutine moves the node x to a level $k \leq j$, then it takes $O(b_x \cdot \alpha^{j+1})$ time.*

Proof (Sketch). Only the edges $(x, y) \in E$ with $\ell(y) \leq j$ get affected as the node x moves to level k. The data structure associated with every other edge remains unchanged. Thus, the total running time of the subroutine is proportional to the number of edges (x, y) with $\ell(y) \leq j$, plus the time for the call to RANDOM-SETTLE(x) or the procedure in Section 2.1.4. The latter quantity is bounded in Lemma 10 and in Section 2.1.4. As we are in Case 2, the former quantity is by definition at most $2b_x \cdot \alpha^{j+1}$. ◀

2.1.3 The subroutine Random-Settle(x)

Suppose that we call RANDOM-SETTLE(x) on a node x at level $\ell(x) = k \geq 0$. From Sections 2.1.1 and 2.1.2, we infer that $2b_x \cdot \alpha^k < |\mathcal{O}_x| \leq 2b_x \cdot \alpha^{k+1}$ at this point in time. Let $\lambda_x = |\mathcal{M}_x|$ be the number of matched edges incident on x just before the call to the subroutine. The subroutine selects $(b_x - \lambda_x)$ edges uniformly at random from $\mathcal{O}_x$ and adds them to $\mathcal{M}$. Thus, at the end of the subroutine we have $|\mathcal{M}_x| = b_x$. The following lemma crucially uses our preferential treatment to FULL-FROM-ABOVE nodes over DEFICIENT nodes.

▶ **Lemma 9.** *If a call to RANDOM-SETTLE(x) matches an edge (x, y) and unmatches an edge (y, z) (Step (04) in Figure 2), then we have $\ell(z) < \ell(x)$ during that call.*

Proof. Consider two possible cases, depending on the state of the node x.

Case 1. The node x is DEFICIENT just before the call to FIX-DIRTY(x) in Figure 1. Then at this stage there is no FULL-FROM-ABOVE node. This holds since the WHILE loop in Figure 1 always picks a FULL-FROM-ABOVE node over a DEFICIENT node. In particular, the node y is not FULL-FROM-ABOVE. Since (y, z) is a matched edge that minimises the value of $\ell(y, z)$, and since y is not FULL-FROM-ABOVE, we must have $\ell(z) \leq \ell(y, z) = \ell(y)$. Finally, note that $(x, y) \in \mathcal{O}_x$, and hence we get: $\ell(z) \leq \ell(y) < \ell(x)$.

Case 2. The node x is FULL-FROM-ABOVE just before the call to FIX-DIRTY(x) in Figure 1. Then at this stage the node x has the highest BASE(x) value among all the FULL-FROM-ABOVE nodes. Suppose that BASE(x) $= j$ at this point in time. Since the subroutine FIX-DIRTY(x) called the subroutine RANDOM-SETTLE(x), we must have been in Case 2 of Section 2.1.1. This means that the node x moves to a level strictly larger than j. Thus, we have $\ell(x) > j$ during the call to RANDOM-SETTLE(x). We now consider two cases, depending on the state of the node y. (a) If the node y is not FULL-FROM-ABOVE during the call to RANDOM-SETTLE(x), then using an argument exactly similar to the one used

in Case 1, we infer that: $\ell(z) \leq \ell(y,z) = \ell(y) < \ell(x)$. This concludes the proof. (b) Else if the node y is FULL-FROM-ABOVE during the call to RANDOM-SETTLE(x), then we claim that BASE(y) $\leq j$ at the same time-instant. This is true since the WHILE loop in Figure 1 picked the node x over y, and so we must have $j = \text{BASE}(x) \geq \text{BASE}(y)$ just before the call to FIX-DIRTY(x). Since (y,z) is an edge in $\mathcal{M}$ that minimises the value $\ell(y,z)$, we get: $\ell(z) \leq \ell(y,z) = \text{BASE}(y) \leq j < \ell(x)$. This concludes the proof. ◀

▶ **Lemma 10.** *A call to the subroutine RANDOM-SETTLE(x) takes $O(b_x \cdot \alpha^{\ell(x)+1})$ time.*

Proof (Sketch). Let $\lambda_x = |\mathcal{M}_x|$ be the number of matched edges incident on x just before the call to the RANDOM-SETTLE(x). The subroutine picks $(b_x - \lambda_x)$ edges uniformly at random from its set of owned edges $\mathcal{O}_x$. This takes $O(|\mathcal{O}_x|)$ time. Further, steps (03) – (04) in each iteration of the WHILE loop in Figure 2 takes $O(1)$ time. Thus, the total time taken by the subroutine is $O(|\mathcal{O}_x| + b_x) = O(b_x \cdot \alpha^{\ell(x)+1})$. The last equality holds since $|\mathcal{O}_x| \leq 2b_x \cdot \alpha^{\ell(x)+1}$ (see the discussion in the beginning of Section 2.1.3). ◀

2.1.4 Handling the nodes in level -1: Maintaining Invariant 2

Every node at level -1 can be in two states: *dead* or *alive*. A node x becomes alive when:

(1) x moves down to level -1 from a higher level. In this case, it scans all its incident edges at level -1 and tries to add them to $\mathcal{M}$ till $|\mathcal{M}_x|$ becomes equal to b_x. If it cannot find sufficient number of edges at level -1 to be FULL, then it remains alive, otherwise it becomes dead. The total computation cost for this step is $O(b_x)$ since x has at most $O(b_x \cdot \alpha^0)$ neighbours in level -1. This cost is charged to Lemma 8.

(2) x is dead at level -1, and it becomes DEFICIENT due to either an unmatching or a deletion of an incident edge. In case (1), x became dead only when it was FULL. Thus, at least ϵb_x edges incident on x have been either deleted from the graph or removed from $\mathcal{M}$. In the former event, we give one unit of credit to x for each edge deletion incident on it. In the latter event, we give one unit of credit to x for every unmatching of an edge incident on it at level $j \geq 0$ from the FIX-DIRTY($\cdot$) procedure that removes this edge from the matching. Thus, node x accumulates at least ϵb_x worth of credit by the time it becomes DEFICIENT. Hence, it can now scan all its incident edges at level -1, as there are $O(b_x)$ such edges.

Finally, if x is not FULL, and an edge (x,y) is added at level -1 where y is also not FULL, then we add the edge (x,y) to the b-matching. This takes $O(1)$ time. A node x at level -1 becomes dead when it is FULL. Once x becomes dead, we wait till it becomes DEFICIENT again. So there is no processing done on x as long as it remains dead. The only non-trivial processing done on x is when it becomes alive, and this takes $O(b_x)$ time. We amortise this cost over the ϵb_x many edge deletions/unmatchings that lead to the change in the state of x.

3 Bounding the Amortised Update Time of Our Algorithm

We fix a sequence of T edge insertions/deletions starting from an empty graph. We show that our algorithm takes $O(T/\epsilon^4)$ time in expectation to handle these T edge insertions/deletions. This implies an amortised update time of $O(1/\epsilon^4)$. Without any loss of generality, we assume that the graph G becomes empty at the end of this sequence of edge insertion/deletions.[1]

[1] Otherwise, we can ourselves delete the existing edges from G one after the other, and get a new sequence of at most $2T$ edge insertions/deletions. We then bound the total update time on this new sequence.

Further, to highlight the main ideas, we only focus on bounding the time spent on handling the nodes at levels $j \geq 0$. From the discussion in Section 2.1.4, it is easy to infer that the amortised time spent on the nodes at level -1 is at most $O(1/\epsilon)$ per update.

Epochs. The notion of an *epoch* will play a key role in our analysis of the amortised update time. We say that a node $x \in V$ *creates an epoch* when it matches a new edge (x, y) during a given iteration of the WHILE loop in RANDOM-SETTLE(v). Note that this is the only way an edge can be added to the b-matching at levels ≥ 0. At most b_v matched edges can be created in RANDOM-SETTLE(v). For the j-th edge selected by v, define the epoch of j-th edge, γ_j to be the contiguous time interval for which this edge remains in the matching. While creating an epoch γ_j, we select an edge (x, y) uniformly at random from $\mathcal{O}_x \setminus \mathcal{M}$ and add this edge to $\mathcal{M}$ (see Step (02) in Figure 2). This random edge (x, y) is called the *candidate edge* of epoch γ_j. We denote the *level of epoch* γ_j by $\ell(\gamma_j)$. This is same as the level of x during the call to RANDOM-SETTLE(x) in Figure 2. We say that an epoch is *destroyed* when its candidate edge is removed from the b-matching $\mathcal{M}$. This happens either when the candidate edge gets deleted from the graph, or when the candidate edge gets unmatched during the course of our algorithm. As the graph G is empty at the end of T edge insertions/deletions, every epoch gets destroyed at some point in time.

Overview of the analysis. For every node $v \in V$ and level $i \in [0, L]$, let $X_{v,i}$ be a random variable that denotes the total number of epochs created by v at level i, as we handle the fixed sequence of T edge insertions/deletions. In Lemma 11, we express the total update time as a function of these random variables $\{X_{v,i}\}$. In Lemma 12, we upper bound the expected value of this function in terms of T. The bound on the expected amortised update time of our algorithm follows from Lemmas 11 and 12 (see Corollary 13).

▶ **Lemma 11.** *It takes* $\sum_{v,i} X_{v,i} \cdot O(\alpha^{i+1}/\epsilon)$ *time to handle* T *edge insertions/deletions.*

Proof (Sketch). The time taken is dominated by the WHILE loop in Figure 1. Hence, it suffices to bound the time spent on the calls to FIX-DIRTY(v) over all $v \in V$. We will charge the total time spent in FIX-DIRTY to the start or an end of an epoch in the following way:

Scenario 1. We have $\ell(v) = i$, v is DEFICIENT, and FIX-DIRTY(v) moves the node v up to some level $j > i$ and calls RANDOM-SETTLE. By Lemma 7 the runtime of FIX-DIRTY(v) is $O(b_v \alpha^{j+1})$. When v comes to level j, it calls RANDOM-SETTLE(v) and starts at least ϵb_v new epochs at level j. So, the computation cost charged per new epoch created is $O(\alpha^{j+1}/\epsilon)$.

Scenario 2. We have $\ell(v) = i$, v is DEFICIENT, and FIX-DIRTY(v) moves v down to some level $j \leq i$. By Lemma 8 the runtime of FIX-DIRTY(v) is $O(b_v \alpha^{i+1})$. Since, v became deficient at level i, at least ϵb_v epochs involving v have been destroyed (since the last time v came to level i, FIX-DIRTY makes it FULL). Note that all such epochs have level $\geq i$. So, the computation cost charged to all epochs destroyed is $O(\alpha^{i+1}/\epsilon)$.

Scenario 3. The hard case is as follows: $\ell(v) = i$, v is FULL-FROM-ABOVE, and FIX-DIRTY(v) moves the node v up to some level $j =$ BASE(v) (or at $j >$BASE(v), Case 2, Section 2.1.1). By Lemma 6 the runtime of FIX-DIRTY(v) is $O(b_v \alpha^{j+1})$. However, it is not necessary that v starts ϵb_v epochs when it arrives at level j. So, we cannot charge this computation cost new epoch of v as no new epochs are created at level j. To overcome this problem, we crucially use the fact that v will eventually become deficient at some higher level. This holds since there are only $\log n$ levels in our partition and v cannot always be

FULL-FROM-ABOVE. Formally, define a phase ϕ of v to be the maximal time interval in which FIX-DIRTY is not called on v. A phase of v is called DEFICIENT if v becomes deficient during the phase, otherwise it is called FULL-FROM-ABOVE. Define a directed graph $\mathcal{H}_v = (\mathcal{U}_v, \mathcal{E}_v)$ where $\mathcal{U}_v$ is the set of all phases of v. For two phases $\phi_1, \phi_2 \in \mathcal{U}_v$, we have $(\phi_1, \phi_2) \in \mathcal{E}_v$ iff ϕ_1 is a FULL-FROM-ABOVE phase and ϕ_2 is the phase that begins just after ϕ_1 ends. If $l(\phi)$ denote the level of v when the phase ϕ starts, then $\ell(\phi_1) < \ell(\phi_2)$. The graph $\mathcal{H}_v$ is a collection of paths and each path ends with a DEFICIENT phase. Let $\Phi = (\phi_1, \phi_2, \ldots, \phi_k)$ be a *maximal* path in $\mathcal{H}_v$. So ϕ_k is a DEFICIENT phase and ϕ_i is a FULL-FROM-ABOVE phase for all $i < k$, and hence $\ell(\phi_1) < \cdots < \ell(\phi_k)$. Let Y_Φ be the total computation cost (due to FIX-DIRTY(v)) in phases in Φ. We get: $Y_\Phi \leq \sum_{i=1}^{k} 2b_v \cdot \alpha^{\ell(\phi_i)+1} = O(b_v \cdot \alpha^{\ell(\phi_k)+1})$. So all the computation cost can be charged to the epochs destroyed during the deficient phase of v, that is $\phi(k)$. When v move to ϕ_k, it is full (or it calls RANDOM-SETTLE(v) to become full. So at least ϵb_v epochs involving v must be destroyed for v to become deficient (All such epochs have level $\geq i$). So we can charge the computation cost Y_ϕ to these ϵb_v epochs and the cost associated with each epoch is $O(\alpha^{i+1}/\epsilon)$.

From Scenario 1,2 and 3, the computation cost charged to each epoch at level i is $O(\alpha^{i+1}/\epsilon)$. So the total time taken by our algorithm is $\sum_{v,i} X_{v,i} O(\alpha^{i+1}/\epsilon)$ ◀

▶ **Lemma 12.** *We have* $\sum_{v,i} E[X_{v,i}].O(\alpha^{i+1}/\epsilon) = O(\alpha^4 T)$.

▶ **Corollary 13.** *The expected amortised update time of our algorithm is* $O(1/\epsilon^4)$.

We now focus on justifying Lemma 12. In Section 3.1, we classify the epochs into three types. Due to space constraints, in Section 3.2 we present a (hand-wavy) intuition behind the proof of Lemma 12. A complete proof appears in the full version of the paper.

3.1 Natural, induced and forced epochs

We say that an epoch is *natural* iff it gets destroyed when its candidate edge is deleted from the graph. If an epoch is not natural, then it gets destroyed when its candidate edge is removed from $\mathcal{M}$ (but does not get deleted from G). This can happen under four possible situations: (1) In Case 2 of Section 2.1.1, when a node x moves up from level j to level $k > j$, we unmatch all its incident edges $(x, y) \in E$ whose other endpoints are at levels $\ell(y) < k$. (2) In Case 1 of Section 2.1.2, when a node x moves up from a level j to a level $k > j$, we unmatch all its incident edges $(x, y) \in E$ whose other endpoints are at levels $\ell(y) < k$. (3) During a call to the subroutine RANDOM-SETTLE(x), we unmatch an edge (y, z) because y gets matched to x. See Step (04) in Figure 1. (4) In Case 2 of Section 2.1.2, when a node x moves down from a level j, we unmatch all its incident edges (x, y) whose other endpoints are at levels $\ell(y) < j$. The epochs whose candidate edges get unmatched under situations (1), (2) and (3) are called *induced* epochs. In contrast, the epochs whose candidate edges get unmatched under situation (4) are called *forced* epochs.

3.2 Intuition behind the proof of Lemma 12: A token based argument

Let $\mathcal{V}$ be the set of all epochs. We assign $\mathcal{T}(\gamma) = \alpha^{\ell(\gamma)+1}$ tokens to every epoch $\gamma \in \mathcal{V}$. Since the computation cost charged to γ (by Lemma 11) is $O(\alpha^{l(\gamma)+1}/\epsilon)$, each token is charged with $O(1/\epsilon)$ amount of computation cost. In Lemma 14, we show that the sum $\sum_{\gamma \in \mathcal{V}} \mathcal{T}(\gamma)$ is at most $O(1/\epsilon)$ times the total number of tokens assigned to the natural epochs,

which is given by the sum $\sum_{\gamma \in \mathcal{N}} \mathcal{T}(\gamma)$. In the paragraph below, we intuitively explain that $E\left[\sum_{\gamma \in \mathcal{N}} \mathcal{T}(\gamma)\right] = O(\alpha \cdot T)$. These observations, taken together, justify Lemma 12.[2]

Suppose that each time an edge (x, y) gets deleted from G, we generate 2 *dollars* and give 1 dollar to each of the endpoints $\{x, y\}$. For the sequence of T edge insertions/deletions in G, the total number of dollars generated is $O(T)$. We will try to convince the reader that a natural epoch γ at level $\ell(\gamma) = i$ receives $\Theta(\alpha^{\ell(\gamma)})$ dollars under this scheme, in expectation. This will give us: $E\left[\sum_{\gamma \in \mathcal{N}} \mathcal{T}(\gamma)\right] = O(\alpha \cdot T)$. Accordingly, suppose that a node x at level $\ell(x) = i$ creates an epoch γ as per Step (02) in Figure 2. From the discussion in the beginning of Section 2.1.3, we get $|\mathcal{O}_x \setminus \mathcal{M}| > b_x \cdot \alpha^i$ at this point in time. We say that each edge $e \in \mathcal{O}_x \setminus \mathcal{M}$ is a *witness* for the epoch γ. Since the sequence of edge insertions/deletions in G is oblivious to the random bits used by the algorithm, in expectation half of these *witness-edges* will get deleted before the candidate edge (x, y) of the epoch γ. Thus, intuitively, if the epoch is natural, then in expectation at least $(b_x/2) \cdot \alpha^i$ of its witness edges get deleted during the lifespan of the epoch. For each of these deletions of the witness edges, the node x receives 1 dollar. Note that each such edge can be a witness to at most b_x epochs of x (since $|\mathcal{M}_x| \leq b_x$). Consider the time-instant when an edge (x, y) gets deleted from G and the node x receives 1 dollar. If we distribute this 1 dollar received by x evenly among all the epochs of x which have (x, y) as a witness, then each of those epochs will receive at least $1/b_x$ dollars. From the preceding discussion, recall that at least $(b_x/2) \cdot \alpha^i$ many witness edges get deleted from G during the lifespan of a natural epoch γ of x at level i. Hence, as promised, we conclude that such an epoch will receive at least $(1/b_x) \cdot (b_x/2) \cdot \alpha^i = \Theta(\alpha^i)$ dollars during its lifespan, in expectation.

▶ **Lemma 14.** *Let $\mathcal{V}$ be the set of all epochs, and let $\mathcal{N} \subseteq \mathcal{V}$ be the set of all natural epochs. Assign $\mathcal{T}(\gamma) = \alpha^{\ell(\gamma)+1}$ tokens to every epoch $\gamma \in \mathcal{V}$. Then $\sum_{\gamma \in \mathcal{V}} \mathcal{T}(\gamma) = O(1/\epsilon) \cdot \sum_{\gamma \in \mathcal{N}} \mathcal{T}(\gamma)$.*

We devote the rest of this section to the proof of Lemma 14. Towards this end, we construct a directed graph $\mathcal{G} = (\mathcal{V}, \mathcal{E})$. To distinguish this from the input graph $G = (V, E)$, we refer to each vertex in $\mathcal{V}$ as a *meta-node* and each edge in $\mathcal{E}$ as a *meta-edge*. The set of meta-nodes $\mathcal{V}$ corresponds to the set of all epochs. The set of meta-edges $\mathcal{E}$ is defined as follows. Each natural epoch has no incoming meta-edge. Each induced or forced epoch has exactly one incoming meta-edge. Thus, without any loss of generality, for every meta-edge $(x, y) \in \mathcal{E}$ directed from x towards y, we say that y is the parent of x and x is a child of y. Note that each meta-node has at most one parent, but it can have multiple children. We will now describe the rules that specify the parent of each induced or forced epoch.

The parent of an induced epoch is specified as follows. Note that an induced epoch is destroyed under three possible scenarios: These are situations (1), (2) and (3) as described in Section 3.1. In situation (3), we define the parent of the induced epoch (with (y, z) as the candidate edge) to be the epoch of x which lead to its destruction (i.e., the epoch with (x, y) as the candidate edge). Both in situation (1) and situation (2), a node x moves up to a higher level, unmatches some incident edges (thereby destroying some induced epochs), and adds some new incident edges to $\mathcal{M}$ (thereby creating some new epochs). As per the descriptions in Sections 2.1.1 and 2.1.2, the node x becomes FULL after these operations. Hence, the number of new epochs created during these steps is at least the number of induced epochs that get destroyed. Accordingly, for each induced epoch γ that gets destroyed, we

[2] This intuitive argument implies that the total expected running time of the algorithm is $O(\alpha \cdot T/\epsilon^2)$. A technically precise argument will lose an extra α multiplicative factor in the running time.

find a unique epoch γ' that gets created, and make γ' the parent of γ. This way of defining parents for the induced epochs ensures the following property.

▶ **Claim 15.** *If an epoch γ' is the parent of an induced epoch γ, then $\ell(\gamma') > \ell(\gamma)$. Further, an epoch γ' can have at most two children that are induced epochs.*

Proof. Let γ' be the parent of an induced epoch γ. Let x and z respectively be the nodes that created the epochs γ' and γ. If γ is destroyed during situation (3) in Section 3.1, then Lemma 9 implies that $\ell(\gamma') = \ell(x) > \ell(y, z) = \ell(\gamma)$ when γ is destroyed. Else if γ is destroyed during situation (1) or (2) in Section 3.1, then the descriptions in Sections 2.1.1, 2.1.2 lead us to the following conclusion: The node x moves up from a level l to some higher level $l' > l$. The epoch γ' is created at level l'. Further, the candidate edge of the epoch γ had both endpoints at a level strictly less than l' when γ was created. Thus, we again get $\ell(\gamma') > \ell(\gamma)$. Finally, an epoch γ' can have at most one child that is an induced epoch which gets destroyed during situation (1) or (2) in Section 3.1, and at most one child that is an induced epoch which gets destroyed during situation (3) in Section 3.1. The claim follows. ◀

We now define the parents of forced epochs. The forced epochs are destroyed during situation (4) in Section 3.1. Thus, we consider the following scenario. A node x is at level $\ell(x) = j \geq 0$, and it becomes DEFICIENT at time t_1 (say). At this stage we have $|\mathcal{M}_x| < (1 - \epsilon) \cdot b_x$. Hence, as the node x moves down, at most $(1 - \epsilon) \cdot b_x$ many forced epochs get destroyed. Let F denote the set of these forced epochs. We have $|F| \leq (1 - \epsilon) \cdot b_x$. Also note that the node x must have been FULL the last time (say t_0) it *settled* at level j after a call to FIX-DIRTY(x). Thus, for situation (4) to occur, the node x must have lost at least ϵb_x many edges in $\mathcal{M}_x$ during the time-interval $[t_0, t_1]$. Let H denote the set of these epochs that were alive at time t_0, were destroyed before time t_1, and whose candidate edges were part of $\mathcal{M}_x$. We have $|H| \geq \epsilon \cdot b_x$.

We claim that $\ell(h) > j$ for every forced epoch $h \in H$. To see why this is true, let (x, y) be the candidate edge for h. Clearly, when h was destroyed the node x stayed put at level j. It follows that since h is a forced epoch, the node y must have created h. The claim is equivalent to the statement that $\ell(y) > j$ when h gets created, and this in turn is equivalent to the statement that $\ell(y) > j$ when h gets destroyed (since $\ell(y)$ does not change during the lifespan of epoch h). To see why the latter statement is true, note that by definition y moves down to a lower level when the epoch h gets destroyed. Thus, as per Case 2 of Section 2.1.2, the node y can unmatch the edge (x, y) only if $\ell(y) > j$ at that time-instant. Hence, we get:

▶ **Property 16.** *Every forced epoch $h \in H$ has level $\ell(h) > j$.*

We now *assign* the epochs in F *evenly* among the epochs in H. To be more specific, after this step, each epoch $h \in H$ gets $|F|/|H| \leq (1 - \epsilon)/\epsilon$ many epochs $f \in F$ *assigned to it*. We are now ready to define the parents of the forced epochs F. Suppose that an epoch $f \in F$ is assigned to an epoch $h \in H$. If $\ell(h) > j$, then h becomes the parent of f. Else if $\ell(h) = j$, then from Property 16 it follows that either h is a natural epoch or h is an induced epoch. In the former event, again h becomes the parent of f. In the latter event, Claim 15 guarantees that h has a parent, say $p(h)$, and this epoch $p(h)$ becomes the parent of f. Claim 15 also implies that $\ell(p(h)) > \ell(h) = j$. Since $\ell(f) = j$, we immediately get the following claim.

▶ **Claim 17.** *Suppose that an epoch γ' is the parent of a forced epoch γ. Then either (1) $\{\ell(\gamma') > \ell(\gamma)\}$ or (2) $\{\gamma'$ is a natural epoch and $\ell(\gamma') \geq \ell(\gamma)\}$.*

Claim 18 holds since: (1) An epoch $h \in H$ gets at most $(1-\epsilon)/\epsilon$ many epochs $f \in F$ assigned to it. (2) An epoch γ can have two children $h, h' \in H$ both of which are induced epochs (see Claim 15), and both h, h' can get at most $(1 - \epsilon)/\epsilon$ many epochs $f \in F$ assigned to them.

▶ **Claim 18.** *An epoch can have at most* $3(1-\epsilon)/\epsilon$ *many forced epochs as its children.*

Proof of Lemma 14. In the meta-graph $\mathcal{G}$, every induced or forced epoch has exactly one incoming edge, and every natural epoch has zero incoming edge. Hence, the meta-graph $\mathcal{G}$ is a collection of rooted directed trees, where the tree-edges are directed away from the roots. The root of each tree is a natural epoch, and any internal node is either an induced epoch or a forced epoch. Let $\mathcal{V}'$ be the set of meta-nodes in any such tree with $r \in \mathcal{V}'$ as its root. To prove the lemma, it suffices to show that $\sum_{\gamma \in \mathcal{V}'} \mathcal{T}(\gamma) = O(1/\epsilon) \cdot \mathcal{T}(r)$.

Only the root r is a natural epoch in $\mathcal{V}'$. Hence, Claims 15, 17, 18 imply that: (1) $\ell(\gamma') > \ell(\gamma)$ whenever an *internal* meta-node $\gamma' \in \mathcal{V}'$ is the parent of $\gamma \in \mathcal{V}'$. (2) Any internal meta-node in $\mathcal{V}'$ has at most $2 + 3(1-\epsilon)/\epsilon \leq 3/\epsilon$ children. (3) The root also has at most $2 + 3(1-\epsilon)/\epsilon \leq 3/\epsilon$ children. Let C be the set of children of the root. Then for all $\gamma \in C$, we have $\ell(r) \geq \ell(\gamma)$.

Consider a meta-node $\gamma \in C$. Let $\mathcal{T}^*(\gamma)$ denote the total number of tokens assigned to the meta-nodes in the subtree rooted at γ. From the above discussions, we get: $\mathcal{T}^*(\gamma) \leq \sum_{j=0}^{\ell(\gamma)} \left(\frac{3}{\epsilon}\right)^{\ell(\gamma)-j} \cdot \alpha^{j+1} = O(\alpha^{\ell(\gamma)+1}) = O(\alpha^{\ell(r)+1})$. The first inequality holds since there are at most $(3/\epsilon)^{\ell(\gamma)-j}$ descendants of γ at level $j \leq \ell(\gamma)$, and each of these descendants get α^{j+1} tokens. The second equality holds since $\alpha = 5/\epsilon >> (3/\epsilon)$. The third equality holds since $\ell(\gamma) \leq \ell(r)$. Hence, the total number of tokens assigned to the tree is given by: $\sum_{\gamma \in \mathcal{V}'} \mathcal{T}(\gamma) = \mathcal{T}(r) + \sum_{\gamma \in C} \mathcal{T}^*(\gamma) \leq \alpha^{\ell(r)+1} + (3/\epsilon) \cdot O(\alpha^{\ell(r)+1}) = O(1/\epsilon) \cdot \mathcal{T}(r)$.

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