

## Exercise 8

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## 1 Sublinear-Time Approximation of Maximum Matching

Consider a graph  $G = (V, E)$ . Recall that a *matching* is a set of edges  $M \subseteq E$  such that no two of the edges in  $M$  share an end-point. A fractional matching is the corresponding natural relaxation, where we assign to each edge  $e \in E$  a value  $x_e \in [0, 1]$  such that the summation of the edge-values in each node is at most 1, that is, for each node  $v \in V$ , we have  $\sum_{e \in E(v)} x_e \leq 1$ , where  $E(v)$  denotes the set of edges incident on  $v$ . We define  $y(v) = \sum_{e \in E(v)} x_e$  as the value of node  $v$  in the given fractional matching. The *size* of a fractional matching is defined as  $\sum_{e \in E} x_e$ , and we have  $\sum_{e \in E} x_e = (\sum_{v \in V} y(v))/2$  (why?). We call a fractional matching *almost-maximal* if for each edge  $e \in E$ , there is one of its endpoints  $v \in e$  such that  $y(v) = \sum_{e' \in E(v)} x_{e'} \geq \frac{1}{1+\epsilon}$ .

### Exercise

- (1a) In the class, we saw that any maximal matching has size at least  $1/2$  of the maximum matching. Prove that the size  $\sum_{e \in E} x_e = (\sum_{v \in V} y(v))/2$  of any almost-maximal fractional matching is at least  $\frac{1}{2(1+\epsilon)}$  of the size of maximum matching.

Consider a maximum matching  $M^*$  and an almost-maximal fractional matching which has value  $x_e$  on each edge  $e$ . We prove that  $\sum_{e \in E} x_e \geq \frac{|M^*|}{2(1+\epsilon)}$ . Consider  $|M^*|$  dollars spread around, where we have put one dollar on each edge  $e$  of the maximum matching  $M^*$ . By the almost-maximality of the fractional matching, each edge  $e$  has at least one endpoint  $v \in e$  such that  $y(v) = \sum_{e' \in E(v)} x_{e'} \geq \frac{1}{1+\epsilon}$ . Make edge  $e$  send its one dollar to one such endpoint  $v$ . This way, each node receives at most one dollar (why?). Now, make node  $v$  split its one dollar among its incident edges  $E(v)$  proportional to the values  $x_{e'}$ . This way, each edge receives at most  $(1+\epsilon)x_{e'}$  dollars from  $v$ . More generally, each edge  $e'$  receives at most  $(1+\epsilon)x_{e'}$  dollars from each of its endpoints and thus overall at most  $2(1+\epsilon)x_{e'}$ . We can conclude that  $\sum_{e' \in E} 2(1+\epsilon)x_{e'} \geq |M^*|$ . In other words,  $\sum_{e \in E} x_e \geq \frac{|M^*|}{2(1+\epsilon)}$ .

Thus, the above item indicates that almost-maximal fractional matchings also provide a reasonable approximation of the size of the maximum matching. But computing an almost-maximal fractional matching is much easier. We next see a LOCAL algorithm for that.

**LOCAL-Algorithm for Almost-Maximal Fractional Matching:** Initially, set  $x_e = 1/\Delta$  for each edge  $e \in E$ . Then, for  $\log_{1+\epsilon} \Delta$  iterations, in each iteration, we do as follows:

- For each vertex  $v$  such that  $y(v) = \sum_{e \in E(v)} x_e \geq \frac{1}{1+\epsilon}$ , we freeze all of its incident edges.
- For each unfrozen edge  $e$ , set  $x_e \leftarrow x_e \cdot (1+\epsilon)$ .

### Exercise

- (1b) Prove that the process always maintains a fractional matching, meaning that we always have  $\sum_{e \in E(v)} x_e \leq 1$ .

Per iteration, we freeze all edges incident on vertices  $v$  whose sum  $y(v)$  has passed  $\frac{1}{1+\epsilon}$  and then we increase unfrozen edges by a  $(1+\epsilon)$  factor. Hence, the value  $y(v)$  can increase to at most  $\frac{1}{1+\epsilon} \cdot (1+\epsilon) = 1$ , but cannot pass that.

- (1c) Prove that at the end, we have an almost-maximal fractional matching, meaning that for each edge  $e \in E$ , there is one of its endpoints  $v \in e$  such that  $\sum_{e \in E(v)} x_e \geq \frac{1}{1+\epsilon}$ .

For each edge  $e$ , either during some it gets frozen because one of its endpoints  $v \in e$  reaches a sum  $y(v) = \sum_{e \in E(v)} x_e \geq \frac{1}{1+\epsilon}$ , or the edge  $e$  gets multiplies by  $(1 + \epsilon)$  in each iteration. The latter means  $x_e$  reaches a value of  $\frac{1}{\Delta} \cdot (1 + \epsilon)^{\log_{1+\epsilon} \Delta} = 1$ . That would imply that even both of the endpoints  $v \in e$  have  $\sum_{e \in E(v)} x_e \geq \frac{1}{1+\epsilon}$ .

Now that we have a simple LOCAL-algorithm for almost-maximal fractional matching, we use it to obtain a centralized algorithm for approximating the maximum matching. To estimate the size of maximum matching, we pick a set  $S$  of  $k = \frac{20\Delta \log 1/\delta}{\epsilon^2}$  nodes at random (sampled with replacement). Here,  $\delta$  is some certainty parameter  $\delta \in [0, 0.25]$ . For each sampled node  $v \in S$ , we run the above LOCAL-algorithm around  $v$ , hence allowing us to learn  $y(v)$ .

### Exercise

- (1d) Define a linear function  $f : \mathbb{R} \rightarrow \mathbb{R}$  such that when applied on the sample average  $\sum_{v \in S} y(v)/|S|$ , the resulting value  $f(\sum_{v \in S} y(v)/|S|)$  is an unbiased estimator of  $\sum_{e \in E} x_e = (\sum_{v \in V} y(v))/2$ . That is,

$$\mathbb{E}_S[f(\sum_{v \in S} y(v)/|S|)] = \sum_{e \in E} x_e.$$

We have  $\mathbb{E}_S[\sum_{v \in S} y(v)/|S|] = \frac{2 \sum_{e \in E} x_e}{n}$  (why?). Hence, it suffices to define  $f(z) = nz/2$ .

- (1e) What is the query complexity of our sublinear-time approximation algorithm?

Per sampled node, we need to simulate the algorithm in its  $(\log_{1+\epsilon} \Delta)$ -hop neighborhood. The size of this neighborhood and thus also the related query complexity is at most  $O(\Delta^{\log_{1+\epsilon} \Delta})$ . Hence, the overall query complexity is  $O(k \Delta^{\log_{1+\epsilon} \Delta}) = O(\Delta^{1+\log_{1+\epsilon} \Delta} \cdot \frac{\log 1/\delta}{\epsilon^2})$ . In terms of dependency on  $\Delta$ , this is much better than the  $2^{O(\Delta)}$  bound that we saw in the class.

- (1f) Prove that the estimator that you defined in (1d) gives a  $(2 + 5\epsilon)$ -approximation of the maximum matching size, with probability at least  $1 - \delta$ .

By (1d), we know that the expectation of our estimator is  $\sum_{e \in E} x_e$ , which we know by (1a) is within a  $2(1 + \epsilon)$  factor of the size of the maximum matching. We next examine how much the random value may deviate from this expectation. Define  $X_i$  to be the random variable that is equal to  $y(s_i)$  where  $s_i$  denotes the  $i^{\text{th}}$  node in our sample set  $S$ . Notice that  $X_i \in [0, 1]$  and moreover,  $\mathbb{E}[X_i] = \frac{2 \sum_{e \in E} x_e}{n}$ . Hence,  $\mu = \mathbb{E}[\sum_{i=1}^k X_k] = \sum_{i=1}^k \mathbb{E}[X_i] = k \frac{2 \sum_{e \in E} x_e}{n}$ . Also notice that  $\sum_{e \in E} x_e \geq \frac{n}{4\Delta}$  (why?) and thus,  $\mu \geq k \frac{n/(2\Delta)}{n} = 20 \frac{\Delta \log 1/\delta}{\epsilon^2} \cdot \frac{1}{2\Delta} = \frac{10 \log 1/\delta}{\epsilon^2}$ . Therefore, by Chernoff bound, the probability that  $X = \sum_{i=1}^k X_k$  deviates by more than a  $(1 + \epsilon)$  factor from its expectation  $\mu$  is at most

$$2e^{-\epsilon^2 \mu / 3} = 2e^{-\frac{10 \log 1/\delta}{3}} \leq \delta.$$

Thus, with probability at least  $1 - \delta$ , we get an expectation within a  $2(1 + \epsilon)(1 + \epsilon) \leq 2 + 5\epsilon$  factor of the maximum matching.