### Lecture 4

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# Plan for today

- 1. More on probability testing.
- 2. Estimate the number of connected components in a graph.

## 1 Testing for monotonicity of a distribution

**<u>Def</u>**: distribution p over domain [n] is "monotone decreasing" if

$$\forall i \in [n-1] : p(i) \ge p(i+1)$$

Goal: design an algorithm such that:

- 1. if p is monotone decreasing output PASS (with probability  $\geq \frac{3}{4}$ )
- 2. if p is  $\epsilon$ -far from monotone decreasing output FAIL (with probability  $\geq \frac{3}{4}$ )

A useful tool - Birge Decomposition: Given any monotone decreasing distribution q and  $\epsilon$ , we decompose the domain [n] into  $l = \Theta(\frac{\log \epsilon n}{\epsilon}) \approx \Theta(\frac{\log n}{\epsilon})$  intervals  $I_1^{\epsilon}, I_2^{\epsilon}, ..., I_l^{\epsilon}$  such that:

$$|I_{k+1}^{\epsilon}| = \left\lceil (1 + \epsilon/2) \cdot |I_k^{\epsilon}| \right\rceil$$

Note: for notation purposes, we disregard  $\epsilon$  and simply denote  $I_k$ .

Define  $\tilde{q}_{\epsilon}$  - "the flattened distribution":  $\tilde{q}_{\epsilon}$  "flattens" each part  $q(I_j)$  of the partition, by distributing uniformly on it's values. Namely:

$$\forall j \in [l], \forall i \in I_j : \tilde{q}_{\epsilon}(i) = \frac{q(I_j)}{|I_j|}$$

Making the original distribution into a "staircase" distribution, where each part of the partition is one stair, and each part keeps it's weight as the original weight it had in q.

### **Important Theorem:**

1. If distribution q is monotone decreasing then  $\|\tilde{q}_{\epsilon} - q\|_1 < \epsilon$ .

2. If distribution q is  $\epsilon$ -close to any monotone decreasing (with respect to  $l_1$  distance) then  $\|\tilde{q}_{\epsilon} - q\|_1 < O(\epsilon).$ 

### Algorithm 1 Testing decreasing monotonicity

- 1: For each part  $I_j$  in the partition: test whether  $q_{|I_j|}$  is close to uniform. If not, output FAIL
- 2:  $w_j \leftarrow$  estimate weights of each partition  $I_j$ .
- 3: Use LP to verify that that w is close to monotone

3. If distribution q is monotone decreasing then for each part  $I_j$  in the partition, we have  $:q_{|I_j|}$  is close to uniform.

Sample analysis: The number of samples required for the above algorithm is

$$\Omega(\frac{\sum \sqrt{I_j}}{\epsilon^2}) \cdot \Theta(\frac{\log n}{\epsilon}) = \Omega(\frac{\sqrt{n} \cdot \log n}{\epsilon^3}) \ .$$

The first term is for testing uniformity in each interval, and the second term is the number of parts in our partition.

### Notes:

1. If at any interval the number of samples is too small approximate by 0.

2. Normally, step 2 is hard, but under the notion that the number of partitions is  $\Theta(\frac{\log n}{\epsilon})$ , the LP is easily solvable.

3. The correctness of this algorithm is also derived from the following observation:

For 2 probability distribution p, q over the same partitions  $I_1^{\epsilon}, I_2^{\epsilon}, ... I_l^{\epsilon}$ , if the conditional distributions hold:  $\forall I_j : \|p_{|I_j} - q_{|I_j}\|_1 < \epsilon$ , then we get:  $\forall I_j : \|p - q\|_1 < \epsilon$ .

4. The first step of Algorithm 1 is checking closeness of each part  $q_{|I_j|}$  to a uniform distribution. The algorithm for that was shown in the previous lecture, and generally, it is not 'tolerant'. Meaning, it might output FAIL for distributions that are  $\epsilon$ -close to uniform.

Luckily, for our possible inputs of  $q_{|I_j}$ , it can be made tolerant enough to keep the correctness of Algorithm 1.

And now for something completely different.

## 2 Estimate the number of connected components in a graph

Given an undirected graph G(V,E), represented as an adjacency-list, and a (relatively small) number dlet us define: n := |V|, m := |E|. we only consider sparse graphs, where d is significantly smaller than n, and  $max_{v \in V} degree(v) \leq d$ .

Generally, sub-linear algorithms over graphs are considered sub-linear time in m, but since it is possible that m=0... particularly in this problem, we will consider the sample complexity to be sub-linear in (m + n).

In this algorithm, we will see that the sample complexity will depend on d, and not on n or m.

**Definition 2.1** A connected component of an undirected graph is a sub-graph in which any two vertices are connected to each other by a path.

We will see an algorithm for estimating the number of connected components in an undirected graph G. our algorithm will return a value y s.t.

$$C - \epsilon n \le y \le C + \epsilon n$$

where C = # connected components. i.e. we get a bound on the distance:  $|y - C| \le \epsilon n$ Notes:

1. There is a lower bound on the sample complexity for this algorithm in terms of  $\epsilon$  and d.

 $2.\epsilon n$  might be very big as to C. Meaning if G is a very big graph (big n), that has little connected components (small C), we might get a big error, related to C.

We note that this does not concern us, since we are interested in the cases where C is big. Namely, we have many connected components in the graph.

We begin with some definitions and observations in order to see how we build such algorithm:

**Definition 2.2** Let v be a vertex in V. We define  $n_v$  as the number of vertices in the connected component to which v belongs. Namely:  $\forall v \in V$ , let  $n_v := \#\{u \in V | \exists path between u and v\}$ 

**Observation 1:**  $\forall$  connected component  $A \subseteq V$ :

$$\Sigma_{v \in A} \frac{1}{n_v} = \Sigma_{v \in A} \frac{1}{|A|} = \frac{|A|}{|A|} = 1 \Rightarrow \Sigma_{v \in V} \frac{1}{n_v} = C$$

(where the rightmost equation comes from the fact that we get 1 over the summation on each and every connoected component A.)

Allegedly: We need  $n^2d$  steps to precisely calculate C. We will now show an approximation that runs in sub-linear time. first, we approximate  $n_v$  and then we approximate the summation itself. We will show that we can estimate  $\sum_{v \in V} \frac{1}{n_v}$  with a small amount of samples, using the standard Chernoff bound. Recall that d is greater than the maximum degree in G. Let us consider d as a constant in the input,  $d \ll n$ .

since the graph is represented as an adjacency list, iterating over all neighbours of a given vertex takes at most d steps. We will estimate  $\sum_{v \in V} \frac{1}{n_v}$  in two steps:

1) estimating  $\frac{1}{n_v}$ 

2) estimating the sum of our values using Chernoff bound.

# Step 1: Estimating $\frac{1}{n_v}$

**Definition 2.3** we define:  $\hat{n}_v := \min\{n_v, \frac{2}{\epsilon}\}$ 

We notice that it means:  $\frac{1}{\hat{n}_v} = max\{\frac{1}{n_v}, \frac{\epsilon}{2}\}$ 

we can assume  $\epsilon$  to be a significantly small number. Namely:  $\epsilon \ll 1$ , Hence,  $\frac{2}{\epsilon} \gg 1$ .

This way, every vertex that belongs to a small connected component, will satisfy:  $\hat{n}_v = n_v$ .

The vertices that belong to large connected component, we can "round down", since the fraction  $\frac{1}{n_v}$  will have a small affect on our summation.

**Definition 2.4** Let us define  $\hat{C}$  as follows:  $\hat{C} = \sum_{v \in V} (\frac{1}{\hat{n}_v})$ 

### Lemma 1:

$$\forall v \, |\frac{1}{\hat{n}_v} - \frac{1}{n_v}| \le \frac{\epsilon}{2}$$

<u>Proof</u>: There are 2 possible cases:

1) if  $n_{v \leq \frac{2}{\epsilon}}$ , then  $\hat{n}_v = n_v \Rightarrow |\frac{1}{\hat{n}_v} - \frac{1}{n_v}| = 0$ 2) else: we have  $n_v > \frac{\epsilon}{2}$ . Therefore: and then  $\hat{n}_v = \frac{2}{\epsilon} \Rightarrow \frac{1}{\hat{n}_v} = \frac{\epsilon}{2}$ ,  $\frac{1}{n_v} < \frac{\epsilon}{2} \Rightarrow |\frac{1}{\hat{n}_v} - \frac{1}{n_v}| = |\frac{\epsilon}{2} - \frac{1}{n_v}| \leq \frac{\epsilon}{2}$ The last inequality stands because  $\frac{1}{n_v}$  is a positive number. Now we will show that:  $|\hat{C} - C| \leq |\sum_{v \in V} (\frac{1}{\hat{n}_v}) - \sum_{v \in V} (\frac{1}{n_v})| \leq \sum_{v \in V} |\frac{1}{\hat{n}_v} - \frac{1}{n_v}| \leq n \cdot \frac{\epsilon}{2} = \frac{\epsilon n}{2}$ where the second inequality comes from pairwise triangle inequality, and the third is true because |V| = n.

And now we have the next consequence:

#### Corollary 1:

$$|\hat{C} - C| \le \frac{\epsilon n}{2}$$

Note that we have constructed the estimation of  $\hat{n}_v$  s.t the estimate of  $|\hat{C} - C|$  can only have half of the error range we had. This gives us room for some additive error in step 2 as well.

(Good question: given i , how can we choose random neighbours? can we check if j is neighbour? Answer: neighbour: running on i's adjacency list O(d) for a neighbour (where d is the bound on the degree in G). check if j is a neighbour: again, running on the list. also O(d))

Now we would like to calculate  $\hat{n}_v$ . How will we do this, and how long will it take?

### **Algorithm 2** calculating $\hat{n}_v$

1: We run a BFS until visiting whole connected component of  $v \underline{\text{or}}$  until we see  $\frac{2}{\epsilon}$  new nodes.

2: we output the number of nodes we visited during that process.

we note that the number of visited nodes is  $= (min\{n_v, \frac{2}{\epsilon}\})$ . <u>Complexity</u>: This is bounded by  $\frac{d*2}{\epsilon}$ . so it's  $O(\frac{d}{\epsilon})$ , since every step of the BFS has time complexity of at most d, and there are at most  $\frac{2}{\epsilon}$  steps of BFS.

So, We can calculate  $\frac{1}{\hat{n}_v}$  in  $O(\frac{d}{\epsilon})$  time for any vertex v.

# Step 2: Estimating $\hat{C}$ .

We start off by describing the algorithm for that calculation

where the choosing of  $r = \frac{b}{\epsilon^3}$ , depends on b, which is a constant that we will choose later on, using Chernoff bound. We will also see (at the proof of Theorem 1), that with high enough probability - that number of samples will suffice.

We notice that the estimation of  $\hat{C}$  is adding us another place for error, since we only estimate it by taking an average over r samples and multiplying it by n.

### **Algorithm 3** estimating $\hat{C}$

- 1: We set  $r := \frac{b}{\epsilon^3}$
- 2: We take **r** samples of  $\hat{n}_v$ .
- 3: choose  $U = \{u_1, ..., u_r\}$  random nodes, uniformly.
- 4:  $\forall u_i \in U$  compute  $\hat{n}_{u_i}$ , using Algorithm 2.
- 5: Sum and output:  $\widetilde{C} = n \cdot \frac{1}{r} (\Sigma_{u_i \in U} \frac{1}{\hat{n}_{u_i}}).$

we will prove later on that most of the times, that estimation is good enough.

A possible problem: summing via the averages could create very rough estimation when dealing with samples that have big variance. for example:

Using that method for (1,2,2,3,4,4,3,2,1,4) will give us a good estimation.

But using that method for  $(0,0,0,0,2^{10000},0,0,0,0)$  will work very badly.

But, since  $\frac{1}{\hat{n}_u} = max\{\frac{1}{n_u}, \frac{\epsilon}{2}\}$ , we get that  $\forall u \in U. \frac{1}{\hat{n}_u} \in \{\frac{1}{n_u}, \frac{\epsilon}{2}\}$ .

Since  $\hat{n}_u \ge 1$  (it is the number of nodes we visit at algorithm 2 - starting at one), we know that  $\frac{1}{\hat{n}_u} \le 1$ , and also by definition  $\hat{n}_u \le \frac{2}{\epsilon}$ , and therefore  $\frac{1}{\hat{n}_u} \ge \frac{\epsilon}{2}$ We finally get that:  $\frac{\epsilon}{2} \le \frac{1}{\hat{n}_u} \le 1$ 

### Theorem 1:

$$Pr[|\widetilde{C} - \widehat{C}| \le \frac{\epsilon}{2} \cdot \widehat{C}] \ge \frac{3}{4}$$

**Proof:** We will use the Chernoff bound:

a little reminder: in general, for  $x_1, ..., x_r$  iid  $x_i \in [0, 1]$  (actually we will even have:  $x_i \in [\frac{\epsilon}{2}, 1]$ ) if we consider  $S = \Sigma x_i, p = E[x_i] = \frac{E[S]}{r}$ , when using Chernoff multiplicative bound, we get:  $Pr[|\frac{S}{r} - p| \ge \delta p] \le Pr[\frac{S}{r} \ge (1 + \delta) \cdot p] + Pr[\frac{S}{r} \le (1 - \delta) \cdot p] \le e^{-\frac{\delta^2 \cdot \mu}{3}} + e^{-\frac{\delta^2 \cdot \mu}{2}} \le 2 \cdot e^{-\frac{\delta^2 \cdot \mu}{3}}$ Where the first inequality comes from union bound. The second from the two multiplicative Chernoff bounds, assuming  $0 < \delta < 1$ , and the third from adding them, taking into account that  $\frac{1}{3} \le \frac{1}{2} \Rightarrow e^{\frac{1}{3}} \le e^{-\frac{1}{3}} \ge e^{-\frac{1}{2}} \Rightarrow e^{-\frac{\delta^2 \cdot \mu}{3}} \ge e^{-\frac{\delta^2 \cdot \mu}{2}}$ 

(An important remark: as long as each sample is chosen uniformly over n nodes, it's o.k if our values (that depend on r) does not seem independent (might as well we have a graph made of cliques of the same size, and all  $n_u$  are equivalent!) - In our case: as long as each  $U_i$  is chosen uniformly and all  $\{U_i\}_{1 \le i \le r}$  are iid.)

So, when using this bound in our case we have:  $p = E_{u \in U}\left[\frac{1}{\hat{n}_u}\right]$ ,  $S = \sum_{i=1}^r \left(\frac{1}{\hat{n}_{u_i}}\right)$ ,  $\delta = \frac{\epsilon}{2}$ We also notice that:  $E_{u \in U}\left[\frac{1}{\hat{n}_u}\right] = E_{u \in V}\left[\frac{1}{\hat{n}_u}\right]$ , since all v's in U are chosen uniformly and independently.

So, we finally get:

$$Pr\left[\left|\frac{1}{r}\sum_{i=1}^{r}\left(\frac{1}{\hat{n}_{u_{i}}}\right) - E_{u\in V}\left[\frac{1}{\hat{n}_{u}}\right]\right| \ge \frac{\epsilon}{2} \cdot E_{u\in V}\left[\frac{1}{\hat{n}_{u}}\right]\right] \le 2 \cdot e^{-\frac{\left(\frac{\epsilon}{2}\right)^{2} \cdot \left(r \cdot E_{u\in V}\left[\frac{1}{\hat{n}_{u}}\right]\right)}{3}}$$

We now want to find such r, so that the above probability would be bounded by  $\frac{1}{4}$ . So we follow this inequality for r:

$$2 \cdot e^{-\frac{\left(\frac{\epsilon}{2}\right)^2 \cdot \left(r \cdot E_{u \in V}\left[\frac{1}{\hat{n}_u}\right]\right)}{3}} \leq \frac{1}{4}$$
$$\Rightarrow e^{-\frac{\left(\frac{\epsilon}{2}\right)^2 \cdot \left(r \cdot E_{u \in V}\left[\frac{1}{\hat{n}_u}\right]\right)}{3}} \leq \frac{1}{8}$$
$$\Rightarrow \log(e^{-\frac{\left(\frac{\epsilon}{2}\right)^2 \cdot \left(r \cdot E_{u \in V}\left[\frac{1}{\hat{n}_u}\right]\right)}{3}}) \leq \log(\frac{1}{8})$$
$$\Rightarrow -\frac{\left(\frac{\epsilon}{2}\right)^2 \cdot \left(r \cdot E_{u \in V}\left[\frac{1}{\hat{n}_u}\right]\right)}{3} \leq -\log(8)$$
$$\Rightarrow \frac{\left(\frac{\epsilon}{2}\right)^2 \cdot \left(r \cdot E_{u \in V}\left[\frac{1}{\hat{n}_u}\right]\right)}{3} \geq \log(8)$$
$$\Rightarrow \frac{\epsilon^2}{4} \cdot \left(r \cdot E_{u \in V}\left[\frac{1}{\hat{n}_u}\right]\right) \geq 3 \cdot \log(8)$$
$$\Rightarrow r \geq \frac{12 \cdot \log(8)}{\epsilon^2} \cdot \frac{1}{E\left[\frac{1}{\hat{n}_{u_i}}\right]}$$

Now, we notice that  $E\left[\frac{1}{\hat{n}_{u_i}}\right] \geq \frac{\epsilon}{2}$ , and therefore  $\frac{1}{E\left[\frac{1}{\hat{n}_{u_i}}\right]} \leq \frac{2}{\epsilon}$ So it's enough that we take r s.t:

$$r \geq \frac{12 \cdot \log(8)}{\epsilon^2} \cdot \frac{2}{\epsilon} \geq \frac{12 \cdot \log(8)}{\epsilon^2} \cdot \frac{1}{E\left[\frac{1}{\hat{n}_{u_i}}\right]}$$

Therefore we get it's enough to take:

$$r_0 := \frac{24 \cdot \log(8)}{\epsilon^3}$$

We also know that  $\frac{1}{r} \Sigma_{i=1}^r \left(\frac{1}{\hat{n}_{u_i}}\right) = \frac{\tilde{C}}{n}$ , and  $E_{u \in V} \left[\frac{1}{\hat{n}_u}\right] = \frac{1}{n} \cdot \Sigma \frac{1}{\hat{n}_u} = \frac{\hat{C}}{n}$ So, finally, taking such  $r_0$  as we defined, we get:

$$Pr\left[|\widetilde{C} - \hat{C}| \ge \frac{\epsilon}{2} \cdot \hat{C}\right] = Pr\left[|\frac{\widetilde{C}}{n} - \frac{\hat{c}}{n}| \ge \frac{\epsilon}{2} \cdot \frac{\hat{c}}{n}\right] \le 2 \cdot e^{-\frac{\left(\frac{\epsilon}{2}\right)^2 \cdot \left(r_0 \cdot E_{u \in V}\left[\frac{1}{\hat{n}_u}\right]\right)}{3}} \le \frac{1}{4}$$

Meaning, we have that:

$$\Pr\left[|\widetilde{C} - \hat{C}| \le \frac{\epsilon}{2} \cdot \hat{C}\right] \ge \frac{3}{4}$$

And thus, we proved Theorem 1!

So, with probability  $\geq \frac{3}{4}$  we also have this inequality holding:

$$\left| \widetilde{C} - \widehat{C} \right| \le \frac{\epsilon}{2} \cdot \widehat{C} \le \frac{\epsilon n}{2}$$
$$\Rightarrow \left| \widetilde{C} - C \right| \le \left| \widetilde{C} - \widehat{C} \right| + \left| \widehat{C} - C \right| \le \frac{\epsilon n}{2} + \frac{\epsilon n}{2} \le \epsilon n$$

when the second inequality in the first row comes from the fact that  $\hat{C} \leq n$ , and the second row's first inequality comes from the triangle inequality.