# Matching Extensions

### 1 Red-Blue Matchings

We are given a bipartite graph G where each is either red or blue. Does G have a perfect matching with exactly k red edges and n-k blue edges? Clearly, assigning cost 1 to red edges and 0 to blue edges and computing a min/max-cost perfect matching does not tell us about intermediate k.

We extend the idea of randomly drawn edge costs from before. For each (i,j) draw  $c_{ij}$  i.u.a.r. from  $\{1,\ldots,2m\}$ . If (i,j) is red, set  $x_{ij}=y\cdot 2^{c_{ij}}$ . If (i,j) is blue, set  $x_{ij}=2^{c_{ij}}$ .

Define B to be the matrix of all  $x_{ij}$ . We have

$$\begin{split} \operatorname{perm}(B) &= \sum_{\sigma \text{ permutations}} \prod_{i} x_{i,\sigma(i)} \\ &= \sum_{\sigma \text{ perfect matchings}} \prod_{i} x_{i,\sigma(i)} \\ &= \sum_{\sigma \text{ permutations}} \prod_{i} \left\{ \begin{aligned} y \cdot 2^{c_{i,\sigma(i)}} & \text{ red edges} \\ 2^{c_{i,\sigma(i)}} & \text{ blue edges} \end{aligned} \end{aligned} \\ &= \sum_{\sigma \text{ perfect matchings}} y^{\operatorname{red}(\sigma)} 2^{\operatorname{cost}(\sigma)} \\ \det(B) &= \sum_{\sigma \text{ perfect matchings}} (-1)^{\operatorname{sgn}(\sigma)} y^{\operatorname{red}(\sigma)} 2^{\operatorname{cost}(\sigma)} \end{split}$$

Combining terms to isolate each occurrence of y,

$$\begin{aligned} \operatorname{perm}(B) &= \sum_{j=0}^{n} y^{j} \sum_{\sigma \text{ perfect matching,red}(\sigma) = j} 2^{\operatorname{cost}(j)} \\ &= \sum_{j=0}^{n} y^{j} \alpha_{j} \\ \det(B) &= \sum_{j=0}^{n} y^{j} \sum_{\sigma \text{ perfect matching,red}(\sigma) = j} (-1)^{\operatorname{sgn}(\sigma)} 2^{\operatorname{cost}(j)} \\ &= \sum_{j=0}^{n} y^{j} \beta_{j} \end{aligned}$$

A perfect matching with exactly k red edges exists iff  $\alpha_k > 0$ . If  $\beta_j \neq 0$  then  $\alpha_j > 0$  for all j (and in particular, j = k). If  $\alpha_j \neq 0$  then  $\beta_j$  may still be 0. But if the min-cost matching among matchings with exactly j red edges is unique, then  $\beta_j \neq 0$ . So, the proofs from last time are easily adapted to show:

**Lemma 1.** The min-cost matching with exactly j red edges is unique with probability at least 1/2.

Assume we are in this case (and repeat for higher probability). It is sufficient to test if  $\beta_k = 0$  (i.e., compute  $\beta_k$ ). Notice that  $\det(B(y)) := p(y)$  is a univariate polynomial in y of degree at most n. We have

$$p(y) = \sum_{j=0}^{n} \beta_j y^j.$$

If we plug in any n+1 distinct values for y and evaluate p(y), we obtain a system of n+1 linear equations for n+1 variables  $\beta_0, \ldots, \beta_n$ . Thus, we can solve the system to get all  $\beta_i$ .

## 2 String Comparisons and Pattern Matching

### 2.1 String Comparisons

We have two strings (could be files, archives, etc) x and y of length n with n very large. They are located in different places. We would like to find out if x = y (with high enough probability) without communicating too much.

The idea is to interpret x and y as n-bit numbers. We choose a random (small) prime p. Send p and x mod p. The recipient computes  $y \mod p$  and compares it to  $x \mod p$ . If they are unequal, then  $x \neq y$ ; otherwise, x = y with TBD probability.

This goes wrong when  $x \neq y$  and  $p = y \mod p$ . Equivalently,  $p \mid (x - y)$ . How many distinct p can divide x - y? Since  $|x - y| \leq 2^n$  (because both are length  $\leq n$ ), |x - y| can have at most n distinct prime factors. So, there are at most n "bad" choices.

If we draw p uniformly from the first  $n^3$  prime numbers, we succeed with probability at least  $1 - n^{-2}$ .

But how big could p get? Equivalently, how large is the  $(n^3)^{th}$  prime? Well, how many primes are  $\leq k$ ? By the prime number theorem, it is about  $k/\ln(k)$ . So the  $(n^3)^{th}$  prime is approximately  $n^3 \log n^3 = \Theta(n^3 \log n)$ . (Whereas  $|x-y| \leq 2^n$ ).

In particular, both p and x mod  $p \le p$  can be encoded in at most

$$\log(\Theta(n^3 \log n)) = \Theta(\log n)$$

bits. This is exponentially better than sending n bits. To boost the probability, we can pick several p independently.

#### 2.2 Pattern Matching

Given a long string x with |x| = n and shorter pattern y with |y| = m, is there an occurrence of y in x? Equivalently, if  $x = x_1, \ldots, x_n$ , find if there is a k with  $x_{k+1} = y_i$  for  $i = 1, \ldots, m$ .

The naive algorithm is  $\mathcal{O}(nm)$  where we check each candidate k and compare all m characters. Suffix tree and related algorithms give  $\mathcal{O}(n+m)$ , but they are not very easy! We'll come up with a much simpler randomized algorithm instead.

Define  $X(k) := x_{k+1}x_{k+2}...x_{k+m}$ . The naive randomized algorithm compares X(k) to y for all k. We can use the previous technique to compute  $X(k) \mod p$  and compare to  $y \mod p$ . However, this takes  $\Theta(m)$  for each computation, so the total is still  $\Theta(mn)$ .

For binary strings, we can do this faster. If we already have X(k), then computing X(k+1) should be faster. Mod p, we can subtract  $x_k 2^{m-1}$  from X(k) to get rid of the first bit, then multiply by two to shift left by one bit. Then we add  $x_{k+m}$ . This is a constant number of operations mod p, so this is  $\Theta(\log p)$  or  $\Theta(1)$  depending on the model.

This leads to our algorithm: choose p randomly from a range TBD, compute  $X(0) \mod p$  and  $y \mod p$ , then compare all  $X(i) \mod p$  to  $y \mod p$  using the efficient update for X(i). If a match is ever indicated, check it in time  $\Theta(m)$ . If verified, return it; otherwise, start over.

For the analysis, we want a union bound over n steps, so we need a false match probability of at most  $1/n^3$ . So, draw p from the first  $n^3$  primes. Therefore,  $\log(p) = \mathcal{O}(\log n)$ . This makes the running time  $\Theta(n \log n + m)$ .

#### 3 Random Walks and Markov Chains

Given an undirected graph, the random walk is at some node in each timestep. It randomly chooses a neighbor to move to next, according to some distribution. Usually, the distribution depends only on the current node, not on previous positions.

In the formal model, a Markov chain is characterized by a state space (in this class, finite)  $\{1,\ldots,n\}$  and transition probabilities  $p_{ij}$  from state i to state j. We write  $P=(p_{ij})_{i,j}$  for the matrix.  $p_{ij}$  is the probability of moving to j given that the walk is currently at i. Thus  $\sum_i p_{ij} = 1$  for all i, so P is stochastic.

The Markov chain starts at time 0 in some state  $x_0$ , and at each discrete timestep t moves to  $x_t$  according to the distribution  $p_{x_{t-1},...}$ . While continuous time versions exist (Brownian motion), we will not study them in this class.

The key Markovian property is

$$\Pr[X_{t+1} = j \mid X_0 = i_0, X_1 = t_1, \dots, X_t = i_t] = \Pr[X_{t+1} = j \mid X_t = i_t].$$

That is, the earlier history does not matter.

A random walk on a graph G is a Markov chain with

$$p_{ij} = \frac{1}{\text{out-degree of } i}.$$

That is, we choose a uniformly random edge. Some natural questions to ask include

- 1. What fraction of time is spent at each node?
- 2. What is the probability of ever making it to j if you are currently at i?
- 3. How long in expectation does it take to get from i to j?
- 4. How long until all nodes have been visited at least once?

A state j for which the probability under (2) is not equal to 1 is called transient – otherwise, it is called persistent. A state can be transient iff the chain can be trapped somewhere else. These are difficult to deal with, so we want to define them away.

**Definition 2.** A Markov chain is irreducible if the graph with edges  $\{(i,j): p_{ij} > 0\}$  is strongly connected.

Another undesirable behavior is oscillations, where we bounce back and forth between nodes. That is, the distribution of states is different at odd or even times t. The periodicity of state i is the gcd of the lengths of all cycles including i. State i is aperiodic iff its periodicity is 1, and periodic otherwise. A Markov chain is aperiodic iff all of its states are aperiodic.