

## Lecture 2 — September 22th, 2014

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## 2.1 Preliminaries: Finding primes

**Note** Although deterministic polynomial-time solutions to the PRIME problem are known (AKS), probabilistic algorithms remain significantly faster (Miller-Rabin's algorithm runs in  $O(\lg(n)^2)$ ).

Fast primality testing algorithms can be used to construct prime-finding algorithms (indeed, no easily computable formula to enumerate prime numbers is known).

### FIND-PRIME

**Input** Integer  $N$

**Output** Prime  $p \in \llbracket N, 2N \rrbracket$

### Algorithm

- Draw  $p$  uniformly from  $\llbracket N, 2N \rrbracket$ .
- Check if  $p$  is prime (e.g. using MILLER-RABIN):
  - If MILLER-RABIN accepts  $p$ , return  $p$ .
  - Otherwise, start over.

**Theorem 2.1 (Chebyshev).** Let  $\pi(x)$  be the number of primes  $\leq x$ . Then  $\pi(x) \geq \frac{x}{2 \ln(x)}$ .

**Theorem 2.2 (The Prime Number Theorem).**  $\pi(x) \underset{x \rightarrow \infty}{\sim} \frac{x}{\ln(x)}$ .

**Corollary 2.3.** The number of primes in  $\llbracket n, 2n \rrbracket$  is  $\Omega\left(\frac{n}{\ln(n)}\right)$ .

**Corollary 2.4.**  $\mathbb{P}_{p \in \llbracket n, 2n \rrbracket} (p \text{ prime}) = \Omega\left(\frac{1}{\ln(n)}\right)$

**Average time complexity** We have  $O(\ln(N))$  iterations by the corollary above; each iteration costs  $O(\ln(N))$  modular additions/multiplications. Hence, the final expected cost is  $O(\ln(N)^2)$ .

**Error** Same as that of MILLER-RABIN.

**Notes** Errors do not accumulate. Also, the number of iterations can be bounded (thus turning this Las Vegas algorithm into a Monte-Carlo one) by failing after a set number of iterations (the probability of returning nothing after  $k$  iterations, or equivalently  $\Theta(k \ln(n))$  operations, would then be  $\frac{1}{2^k}$ ).

## 2.2 Polynomial identity testing

### 2.2.1 Problem definition

#### POLYNOMIAL-IDENTITY-TESTING (PIT)

**Input**  $Q$  and  $R$ , two  $n$ -variables polynomials of degree  $\leq d$ .

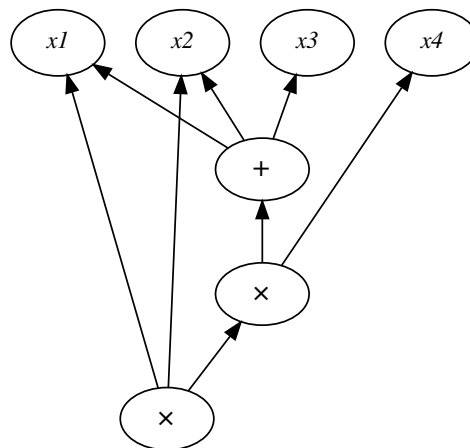
**Output** ACCEPT iff  $Q = R$ .

**Notes** Expanding  $P$  and  $Q$  and comparing individual coefficients takes exponential time in the size of their representation – in other words, compact representations exist that allow for fast evaluation of polynomials whose expanded form contains an exponential number of coefficients.

In the **black-box** model nothing is known about  $P$  and  $Q$ , and the only available operation is  $x \mapsto P(x), Q(x)$ . This single operation is assumed to be fast.

**Example 1: Determinant** Let  $Q = \prod_{1 \leq i < j \leq n} (X_i - X_j)$  and  $R = \det(X_i^j)$ . Then  $Q = R$ , evaluating  $Q$  and  $R$  takes linear time in  $n$ , and expanding  $Q$  and  $R$  takes exponential time in  $n$ .

**Example 2: Arithmetic circuits** Arithmetic circuits are a tree-based representation of polynomial factorizations.



**Figure 2.1.**  $x_1x_2x_4(x_1 + x_2 + x_3)$

**State of the art** Deterministic solutions for the PIT problems are known for polynomials represented as arithmetic circuits of depth  $\leq 2$ . Partial results were also obtained for multi-linear polynomials of depths 3, 4.

(Additional note: depth 4 is the most important one; deterministically solving PIT for arithmetic circuits of depth 4 would represent a significant leap forward for complexity theory.)

**Lemma 2.5 (Schwartz-Zippel).** Let  $F$  denote an arbitrary field, and  $S$  denote a finite subset of  $F$ . Then for any non-zero polynomial  $T(X_1, \dots, X_n)$  of degree  $d$ ,

$$\mathbb{P}_{a_1, \dots, a_n \in S} (T(a_1, \dots, a_n) = 0) \leq \frac{d}{|S|}$$

**Proof (by induction):** If  $n = 1$ , then  $T$  has at most  $d$  roots, and  $\mathbb{P}_{a \in S} (T(a) = 0) \leq \frac{d}{|S|}$ . If  $n > 1$ , expanding  $T$  by its first variable yields  $T = \sum_i X_1^i T_i(X_2, \dots, X_n)$ . Let  $j$  be the degree of  $T$  relative to  $X_1$  – that is, the highest  $i$  such that  $T_i \neq 0$ . Then

$$\begin{aligned} \mathbb{P}_{a_1, \dots, a_n \in S} (T(a_1, \dots, a_n) = 0) &= \mathbb{P}_{a_1, \dots, a_n \in S} (T(a_1, \dots, a_n) = 0 \text{ and } T_j(a_2, \dots, a_n) = 0) \\ &\quad + \mathbb{P}_{a_1, \dots, a_n \in S} (T(a_1, \dots, a_n) = 0 \text{ and } T_j(a_2, \dots, a_n) \neq 0) \end{aligned}$$

Noting that  $T_j$  is a  $n - 1$  variables polynomial of degree  $d' = d - j$  and applying the induction hypothesis yields  $\mathbb{P}_{a_1, \dots, a_n \in S} (T_j(a_2, \dots, a_n) = 0) \leq \frac{d-j}{|S|}$ , which implies that

$$\mathbb{P}_{a_1, \dots, a_n \in S} (T(a_1, \dots, a_n) = 0 \text{ and } T_j(a_2, \dots, a_n) = 0) \leq \frac{d-j}{|S|}$$

To bound the second term, introduce  $a_2, \dots, a_n$  such that  $T_j(a_2, \dots, a_n) \neq 0$ . The strong induction hypothesis applied to  $T(X_1, a_2, \dots, a_n)$  (a single-variable polynomial of degree  $j$ ) yields  $\mathbb{P}_{a_1 \in S} (T(a_1, \dots, a_n) = 0) \leq \frac{j}{|S|}$ . In other words,

$$\mathbb{P}_{a_1, \dots, a_n \in S} \left( \underbrace{T(a_1, \dots, a_n) = 0}_E \mid \underbrace{T_j(a_2, \dots, a_n) = 0}_F \right) \leq \frac{j}{|S|}$$

Finally, note that

$$\begin{aligned} \mathbb{P}_{a_1, \dots, a_n \in S} (T(a_1, \dots, a_n) = 0 \text{ and } T_j(a_2, \dots, a_n) \neq 0) &= \mathbb{P}(E \cup F) \\ &= \mathbb{P}(E \mid F) \mathbb{P}(F) \\ &\leq \mathbb{P}(E \mid F) \\ &\leq \frac{j}{|S|} \end{aligned}$$

Combining both results yields the stated inequality:

$$\mathbb{P}_{a_1, \dots, a_n \in S} (T(a_1, \dots, a_n) = 0) \leq \frac{d}{|S|}$$

□

### Algorithm

- Draw  $\vec{a} = a_1, \dots, a_n$  randomly from  $S = \llbracket 1, 2d + 1 \rrbracket$
- Accept iff  $P(a_1, \dots, a_n) = Q(a_1, \dots, a_n)$

**Time complexity** Two polynomial evaluations.

**Error**

- One sided
- True-biased

If  $P \neq Q$ , then by Schwartz-Zippel's lemma  $\mathbb{P}(ACCEPT) = \mathbb{P}_{\vec{a} \in S} (P(\vec{a}) = Q(\vec{a})) = \mathbb{P}_{\vec{a} \in S} \left( \underbrace{P(\vec{a}) - Q(\vec{a})}_{T(\vec{a})} = 0 \right) \leq \frac{d}{|S|} = \frac{d}{2d+1} < \frac{1}{2}$ .

**Notes** In practice, evaluating  $P$  and  $Q$  can yield extremely large values. To circumvent this problem, all calculations are generally made modulo a large prime value  $p$ . Carefully choosing this value is crucial to ensure that  $P = Q \pmod{p}$  is indeed equivalent to  $P = Q$ . Denoting the largest coefficient of  $P$  and  $Q$  as  $M$ ,  $p$  can be obtained by choosing a prime value larger than twice the maximum of  $d$  and  $M$ .

## 2.2.2 Application to Bipartite perfect matching

### BIPARTITE-PERFECT-MATCHING (BPM)

**Input** Balanced bipartite graph  $G = (E, U \sqcup V)$ , with  $|U| = |V| = n$ .

**Output** ACCEPT iff a perfect matching exists in  $E$ , i.e.  $E$  contains  $n$  disjoint edges.

**Note** A deterministic  $O(\sqrt{|U| + |V|} \cdot |E|) = O(n^{2.5})$  time solution yielding such a perfect matching if it exists is known (Hopcroft-Craft). Probabilistic algorithms by Lovasz (1979) achieve a time complexity for the decision problem equal to that of the calculation of a single  $n \times n$  determinant modulo  $p \in \llbracket n, 2n \rrbracket$ . A 1987 extension by Mulmuley, U. Vazirani, and V. Vazirani gives a probabilistic estimate of the largest such matching in any general graph, in  $O(1)$  matrix inversions time.

**Note** The calculation of a determinant can be reduced to a matrix multiplication problem.

**Adjacency matrices** Identify  $u$  and  $V$  with  $\llbracket 1, n \rrbracket$ , and define the bi-adjacency matrix  $A$  as

$$A_{i,j} = \begin{cases} 1 & \text{if } (i,j) \in E \\ 0 & \text{if } (i,j) \notin E \end{cases}$$

Expanding  $\det(A)$  yields  $\det(A) = \sum_{\sigma \in \mathfrak{S}_n} (-1)^{\text{sgn}(\sigma)} \prod_i A_{i,\sigma(i)}$ , and  $\prod_i A_{i,\sigma(i)}$  is non-zero iff  $\sigma$  represents a perfect matching in  $G$ . Hence if  $\det(A) \neq 0$  then there exists at least one perfect matching. The converse, unfortunately, does not hold due to the  $(-1)^{\text{sgn}(\sigma)}$  term.

**Note** The permanent of  $A$ , defined as  $\text{perm}(A) = \sum_{\sigma \in \mathfrak{S}_n} \prod_i A_{i,\sigma(i)}$ , exactly equals the number of BPM in  $G$ , but computing it is a #P-complete problem; the fastest known deterministic solution (Ryser's formula) has  $O(2^n n)$  time complexity. The fastest known approximation (Jerrum, Sinclair and Vigoda) still requires  $O(n^{10})$  time.

**Tutte matrix** Since computing the determinant of  $A$  is not sufficient, we introduce the Tutte matrix  $T$  of  $G$  as the  $n \times n$  matrix

$$T_{i,j} = \begin{cases} X_{i,j} & \text{if } (i,j) \in E \\ 0 & \text{if } (i,j) \notin E \end{cases}$$

**Theorem 2.6.**  $\det(T)$  is a  $|E|$ -variables polynomial of  $\mathbf{Z}_2^n[X]$  whose degree  $d$  is  $\leq n$ , and  $\det(T) \neq 0 \iff G$  has a BPM.

**Proof:** If no BPM exists, then the determinant is null. Conversely, if a BPM exist, then the determinant is non-null. Indeed, each non-zero  $\prod_i \delta_{(i,\sigma(i)) \in E} X_{i,\sigma(i)}$  monomial in the expansion of  $\det(T)$  matches a single permutation, and is thus distinct of all other monomials in the expanded  $\det(T)$  polynomial.  $\square$

Since the elements of  $T$  are polynomials, expanding  $\det(T)$  is extremely costly. On the other hand, since  $\forall x, \det(T)(x) = \det(T(x))$ , evaluating  $\det(T)$  in a single point is relatively cheap.

### Algorithm

- Pick a prime number  $p \in \llbracket n^2, 2n^2 \rrbracket$ .
- Draw  $|E|$  random elements  $(a_i)$  from  $\llbracket 1, p-1 \rrbracket$ .
- Accept iff  $\det(T(a)) \neq 0 \pmod p$ , where  $T$  is the Tutte matrix of  $G$ .

### Error

- One sided
- False-biased (If the algorithm accepts, then the existence of a BPM is guaranteed)

The probability of incorrectly rejecting is exactly  $\mathbb{P}(\det(T)(a) = 0 \mid \det(T) \neq 0)$ , which by Schwartz-Zippel's lemma is  $\leq \frac{d}{|S|} \leq \frac{n}{n^2} = \frac{1}{n}$ .

**Time complexity** Equal to that of computing an  $n \times n$  determinant ( $O(n^{2.3727})$  using Coppersmith-Winograd algorithms).

## 2.3 Exercises

### 2.3.1 Fingerprints

**FINGERPRINT** Let A and B denote two players.

**First player's input**  $n$ -bits sequence  $u \in \{0, 1\}^n$ .

**Second player's input**  $n$ -bits sequence  $v \in \{0, 1\}^n$ .

**Output** ACCEPT iff  $u = v$ .

**Complexity** Number of bits exchanged.

**Naive solution**

- A sends  $u$  to B.
- B accepts iff  $u = v$ .

**Complexity**  $n$  bits.

**Hash functions** Vectors of  $\mathbf{Z}_2^n$  are mapped to elements of  $\mathbf{Z}_2[X_1, \dots, X_n]$  through the hash function  $H : (a_i) \mapsto \sum_{0 \leq i < n} a_{i+1} X^i$  (or  $\tilde{H} : (a_i) \mapsto \sum_{0 \leq i < n} a_{n-i} X^i$ ). These functions are such that  $H(u) = H(v) \iff u = v$ .

**Algorithm**

- A picks a prime number  $p \in \llbracket n^2, 2n^2 \rrbracket$ .
- A picks a random number  $a \in \llbracket 1, n-1 \rrbracket$ .
- A sends  $(p, a, H(u)(a) \bmod p)$  to B.
- B accepts iff  $H(v)(a) = H(u)(a) \bmod p$ .

**Error**

- One-sided
- True-biased

If  $u \neq v$ , then B accepts with probability  $\leq \frac{1}{n}$ .

**Complexity**  $6 \lg(n) + o(1)$  bits.

**Time complexity**  $n$  modular additions and multiplications for both A and B.

**Note** This algorithm is insecure: it is vulnerable to collision-based attacks.

**2.3.2 Pattern-matching****PATTERN-MATCHING**

**Input** Word  $w \in \mathbf{Z}_2^n$ , pattern  $p \in \mathbf{Z}_2^k$ .  $k \leq n$ .

**Output** Positions where  $p$  occurs in  $w$ :  $\{i \mid p = w[i : i+k-1]\}$ .

**Note** A naive deterministic algorithm (for each index  $i \in \llbracket 1, n-k+1 \rrbracket$  in  $w$ , check whether  $p = w[i : i+k-1]$ ) runs in  $O(nk)$  time. Many efficient, deterministic, linear-time solutions are known (Rabin–Karp, Knuth–Morris–Pratt, Boyer–Moore, etc.), but all are tricky to implement. Probabilistic algorithms, on the other hand, achieve similar performance and are very easy to implement.

**Note** The nature of our hash functions allows for easy calculation of checksums of overlapping subwords. Recall that  $\tilde{H} : (a_j) \mapsto \sum_{0 \leq j < n} a_{n-j} X^j$ , and assume that  $h_i(a) = \tilde{H}(w[i : i+k-1])(a) = \sum_{0 \leq j < k} w_{i-1+(k-j)} a^j$  is known. Then  $h_{i+1}(a)$  can be derived in  $O(1)$  from  $h_i$ . Indeed,

$$\begin{aligned} h_{i+1} &= \tilde{H}(w[i+1 : i+k]) \\ &= \sum_{0 \leq j < k} w_{i+(k-j)} X^j \\ &= \sum_{1 \leq j < k} w_{i+(k-j)} X^j + w_{i+k} \\ &= X \sum_{0 \leq j < k-1} w_{i-1+(k-j)} X^j + w_{i+k} \\ &= X(h_i - w_i X^{k-1}) + w_{i+k} \end{aligned}$$

Evaluating in  $a$  yields  $h_{i+1}(a) = w_{i+k} + a(h_i(a) - w_i a^{k-1})$ .

**Algorithm** As usual, all calculations are run modulo a large enough prime value  $q$ . For each index  $i$ , we decide whether  $w[i : i+k-1]$  matches the pattern  $p$  by comparing  $h_i(a)$  to  $\tilde{H}(p)(a)$ , for randomly sampled values of  $a$ .

- Pick a prime number  $q \in \llbracket n^3, 2n^3 \rrbracket$ .
- Draw  $a$  randomly from  $\llbracket 0, q-1 \rrbracket$ .
- Compute  $h_p = \tilde{H}(p)(a)$ .
- Compute  $h = \tilde{H}(w[1 : k])$ .
- For  $i \in \llbracket 1, n-k+1 \rrbracket$ 
  - If  $h = h_p$ , then append  $i$  to the list of accepted indices.
  - If  $i \neq n-k+1$ , then update  $h \leftarrow w_{i+k} + a(h - w_i a^{k-1})$ .

**Time complexity**  $O(n)$  modular additions/multiplications.

### Error

- One-sided
- True-biased

Errors consist in returning extraneous indices. For each non-matching index  $i$ ,

$$\begin{aligned} \mathbb{P}(i \in \text{returned-values}) &= \mathbb{P}(h_i(a) = \tilde{H}(p)(a) \mid h_i \neq \tilde{H}(p)) \\ &\leq \frac{k}{p} \leq \frac{k}{n^3} \leq \frac{1}{n^2} \end{aligned}$$

Hence the union bound yields

$$\mathbb{P}(\text{incorrect output}) = \mathbb{P}(\exists i \in \text{returned-values} \mid p \neq w[i : i+k-1]) \leq n \cdot \frac{1}{n^3} \leq \frac{1}{n^2}$$

**Note** Instead of choosing large prime numbers, one can reduce the probability of error by computing checksums for multiple different  $a$ .

### 2.3.3 Associativity testing

**ASSOCIATIVE**  $S = \llbracket 1, n \rrbracket$

**Input**  $\circ : S \times S \rightarrow S$ .

**Output** ACCEPT iff  $\circ$  is associative.

**Complexity** Number of operations involving  $\circ$ .

**Naive solution** Checking all possible triples  $(i, j, k) \in S^3$  requires  $2n^3$  comparisons, and (assuming proper memoisation)  $n^2$  evaluations of  $\circ$ .

**Notes** The number of witnesses of the non-associativity of an arbitrary law  $\circ$  may be very small. As an example, consider defining  $i \circ j = 3$  for all  $i, j$  except  $1 \circ 2 = 1$ . Then for all  $\forall(a, b, c) \neq (1, 2, 2)$ ,  $a \circ (b \circ c) = 3 = (a \circ b) \circ c$ , but  $(1 \circ 2) \circ 2 = 1 \neq 3 = 1 \circ (2 \circ 2)$ . In this case there exists a single witness  $(1, 2, 2)$  of the non-associativity of  $\circ$ . The following sections are hence dedicated to expanding the search space to increase the relative frequency of witnesses.

**Extension of the search space** Let  $S(p) = (\mathbf{Z}_p)^n$ , and let  $(e_1, \dots, e_n)$  denote a basis of  $S(p)$ . Define the bilinear  $\bullet$  operation over  $S(p)$  by taking  $e_i \bullet e_j = e_{i \circ j}$  and extending it to  $S(p)$ . Finally, note that if  $(A_i)_i$  denotes the coefficients of  $A$  in the  $(e_i)_i$  basis, then  $A \bullet B = \sum_{i,j} A_i B_j e_{i \circ j}$ .

**Lemma 2.7.**  $\bullet$  is associative iff.  $\circ$  is.

**Proof:** Assume  $\circ$  is associative. Then  $\forall(i, j, k), (e_i \bullet e_j) \bullet e_k = e_{(i \circ j) \circ k} = e_{i \circ (j \circ k)} = e_i \bullet (e_j \bullet e_k)$ .

Conversely, assume  $\bullet$  is associative. Then  $\forall(i, j, k), e_{(i \circ j) \circ k} = (e_i \bullet e_j) \bullet e_k = e_i \bullet (e_j \bullet e_k) = e_{i \circ (j \circ k)}$ , and hence  $(i \circ j) \circ k = i \circ (j \circ k)$ .  $\square$

**Lemma 2.8.** For all  $(A, B, C) \in S(p)$ ,  $(A \bullet B) \bullet C$  is a third-degree polynomial in the coefficients of  $A, B, C$ .

**Proof:** Explicit expansion yields  $(A \bullet B) \bullet C = \sum_{i,j,k} A_i B_j C_k e_{(i \circ j) \circ k}$ .  $\square$

**Lemma 2.9.** Assume that  $p = 7$  and that  $\circ$  is not associative.

Then  $\mathbb{P}_{A,B,C \in S} ((A \bullet B) \bullet C = A \bullet (B \bullet C)) \leq \frac{3}{7}$ .

**Proof:** Given that  $\circ$  is not associative, there exists a 3-tuple  $(A, B, C) \in S(p)^3$  such that  $(A \bullet B) \bullet C \neq A \bullet (B \bullet C)$ . In other words, the third-degree polynomial  $(A \bullet B) \bullet C - A \bullet (B \bullet C)$  in the  $A_i, B_j, C_k$  coefficients is not null. Hence (Schwartz-Zippel)  $\mathbb{P}_{A,B,C \in S} ((A \bullet B) \bullet C = A \bullet (B \bullet C)) \leq \frac{d}{\#S(p)} = \frac{3}{7}$ .  $\square$



**Algorithm**

- Draw  $A, B, C$  at random from  $S(7)$ .
- Compute  $AB = A \circ B$ ,  
 $BC = B \circ C$ ,  
 $AB\_C = AB \circ C$ ,  
 $A\_BC = A \circ BC$ .
- Accept iff.  $AB\_C = A\_BC$ .

**Complexity**  $n^2$  calls are required to build the full multiplication table of  $\circ$ .

**Time complexity** Each of the four subsequent calculations require  $O(n^2)$  modular additions and multiplications, bringing the total time complexity to  $O(n^2)$ .

**Error**

- One-sided
- True-biased

If  $\circ$  is not associative, then (by lemma 2.9)  $\mathbb{P}(ACCEPT) \leq \frac{3}{7}$ .