- 教材讨论
  - -JH第5章第3节第4小节

# 问题1: Neq-Pol

Describing the idea of following algorithm.

#### Algorithm 5.3.4.4. NEQ-POL

```
Input: Two polynomials p_1(x_1,\ldots,x_m) and p_2(x_1,\ldots,x_m) over \mathbb{Z}_n with at most degree d, where n is a prime and n>2dm. Step 1: Choose uniformly a_1,a_2,\ldots,a_m\in\mathbb{Z}_n at random. Step 2: Evaluate I:=p_1(a_1,a_2,\ldots,a_m)-p_2(a_1,a_2,\ldots,a_m). Step 3: if I\neq 0 then \mathrm{output}(p_1\not\equiv p_2) {accept} else \mathrm{output}(p_1\equiv p_2) {reject}.
```

What does step 2 means? Why?

# Neq-Pol is one-sided error Monte Carlo algorithm

 Explain the basic idea of proving the following theorem.

**Theorem 5.3.4.5.** Algorithm Neq-Pol is a polynomial time one-sided-error Monte Carlo algorithm that decides the nonequivalence of two polynomials.

- If 
$$p_1 = p_2$$
  $Prob(Neq-Pol rejects (p_1, p_2)) = 1$ .  
- If  $p_1 \neq p_2$  
$$Prob(Neq-Pol accepts (p_1, p_2)) = Prob(p_1(a_1, \dots, a_m) - p_2(a_1, \dots, a_m) \neq 0) \geq 1 - \frac{m \cdot d}{n} \geq \frac{1}{2}$$
.

# 问题2: Fingerprinting

 Why is NEQ-POL an application of fingerprinting?

we test whether the fingerprint  $p_1(a_1, \ldots, a_n)$  of  $p_1$  is identical to the fingerprint  $p_2(a_1, \ldots, a_2)$  of  $p_2$  for random  $a_1, \ldots, a_n$ .

 What is the concrete meanings of error Prob. for NEQ-POL?

## 问题3: EQ-1BPs

#### What is 1BPs?

Equivalence problem for one-time-only branching programs.

The equivalence problem for one-time-only branching programs, Eq-1BP, is to decide, for two given one-time-only branching programs  $B_1$  and  $B_2$ , whether  $B_1$  and  $B_2$  represent the same Boolean function. One can represent a branching program in a similar way as a directed weighted graph<sup>28</sup> and so we omit the formal description of branching program representation.<sup>29</sup>

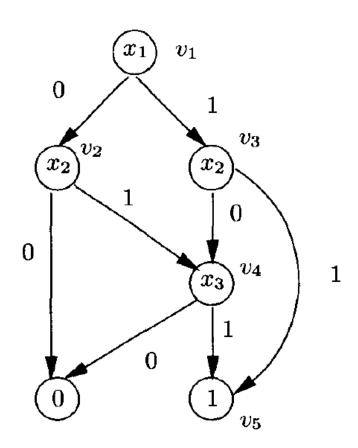
#### EQ-1BP

Input: One-time-only branching program  $B_1$  and  $B_2$  over a set of Boolean variables  $X = \{x_1, x_2, x_3, \ldots\}$ .

Output: "yes" if  $B_1$  and  $B_2$  are equivalent (represent the same Boolean function),

"no" otherwise.

### Constructing a Polynomial for a 1BP



### The Properties

**Observation 5.3.4.7.** For every 1BP A over the set of variables  $\{x_1, x_2, \ldots, x_m\}$ ,

- (i)  $p_A(x_1, \ldots, x_m)$  is a polynomial of degree at most 1 for every variable,
- (ii)  $p_A(a_1,\ldots,a_m)=A(a_1,\ldots,a_m)$  for every Boolean input  $(a_1,\ldots,a_m)\in\{0,1\}^m$ .

**Lemma 5.3.4.8.** For every two 1BPs A and B, A and B are equivalent if and only if  $p_A$  and  $p_B$  are identical.

### Algorithm: NEQ-1BP

What is the idea of the following algorithm?

#### **Algorithm 5.3.4.9.** NEQ-1BP

```
Input: Two 1BPs A and B over the set of variables \{x_1, x_2, \ldots, x_m\}, m \in \mathbb{N}.
```

Step 1: Construct the polynomials  $p_A$  and  $p_B$ .

Step 2: Apply the algorithm NEQ-POL on  $p_A(x_1,\ldots,x_m)$  and  $p_B(x_1,\ldots,x_m)$ 

 $x_m$ ) over some  $\mathbb{Z}_n$ , where n is a prime that is larger than 2m.

Output: The output of NEQ-POL.

What is the essential strategy of NEQ-1BP?