

COMPSCI 501: Formal Language Theory

Lecture 34: Probabilistic Algorithms

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Outline

- ▶ Models of probabilistic acceptance
- ▶ Primality Testing
- ▶ Branching Programs
- ▶ Polynomials for Evaluation

Probabilistic Turing Machines

Def.: Nondeterministic TM with

- ▶ two branches at each nondeterminism step
- ▶ equal probabilities for both (coin flip) $\implies \Pr[b] = 2^{-k}$ for branch b with k coin flips

$$\text{Acceptance: } \Pr[M \text{ accepts } w] = \sum_{\text{accepting } b} \Pr[b]$$

$$\Pr[M \text{ rejects } w] = 1 - \Pr[M \text{ accepts } w]$$

M **decides language A with error probability ϵ** if

- ▶ $w \in A \implies \Pr[M \text{ accepts } w] \geq 1 - \epsilon$
- ▶ $w \notin A \implies \Pr[M \text{ rejects } w] \geq 1 - \epsilon$

BPP (Bounded Error Probabilistic Polynomial Time)

BPP is the class of languages decided by probabilistic *polynomial* Turing machines with error probability $1/3$

Choice of $1/3$ is arbitrary. Anything in $(0, 1/2)$ works. We can make the error probability arbitrarily small.

Amplification Lemma. For any polynomial $p(n)$, a PPTM M_1 with error probability $< 1/2$ has an equivalent M_2 with err. prob. $2^{-p(n)}$

= arbitrary polynomial exponent, still in polynomial time

How: M_2 simulates M_1 $2k$ times, takes majority decision.

Max. error prob. on one run: k right, k wrong, $\epsilon^k(1-\epsilon)^k$.
Over all 2^{2k} result sequences: $2^{2k} \epsilon^k(1-\epsilon)^k = (4\epsilon(1-\epsilon))^k$.

We want $(4\epsilon(1-\epsilon))^k \leq 2^{-p(n)}$. $\implies k = -p(n)/\log(4\epsilon(1-\epsilon))$

Primality Testing

Fermat's Theorem: If $a \in \mathbb{Z}_p^+$ (p prime), then $a^{p-1} \equiv 1 \pmod{p}$

Corollary: if $a < n$, and $a^{n-1} \not\equiv 1 \pmod{n}$, then n is not prime.

Fermat test: n passes test at a if $a^{n-1} \equiv 1 \pmod{n}$

Pseudoprime: n passes Fermat test for all $0 < a < n$. There are non-prime numbers that pass all tests (Carmichael numbers)

Can prove: if p is not pseudoprime, it fails at least half the tests.
 \implies run for k values a_i , error bound 2^{-k} .

Miller-Rabin Primality Testing: Square Roots of 1

1 only has square roots -1 and 1 modulo any prime p .

p prime candidate \implies odd \implies check $a^{\frac{p-1}{2}}$

if $\neq \pm 1$, p is not prime.

If 1, can keep dividing exponent as long as even.

Write $p-1 = s \cdot 2^t$.

Choose $a < p$.

If $a^{p-1} \not\equiv 1 \pmod{p}$, *reject* (p not prime)

Compute sequence $a^{s2^0}, a^{s2^1}, a^{s2^t} \pmod{p}$

If some element $\neq 1$, and last such element $\neq -1$, *reject*

Repeat for k different values $a \implies$ error probability $\leq 2^{-k}$

Theorem: PRIMES is in BPP.

Primality Testing: Proof

- ▶ If number is rejected, it's composite
 - ▶ by Fermat's theorem, or
 - ▶ $b^2 \equiv 1 \pmod{p} \implies (b+1)(b-1) = cp$
- ▶ If number is accepted, very likely prime

Each composite non-witness has a unique corresponding witness

The Class RP

Our test had **one-sided error**

- ▶ *accept* means likely prime
- ▶ *reject* means surely composite

Def. **RP** is the class of languages decided by PPTM, where strings in the language are accepted with probability $\geq 1/2$, and strings not in the language are rejected with probability 1.

We showed $COMPOSITES \in RP$

Another primality test (Adleman-Huang) always rejects composites, and accepts primes with $\Pr \geq 1/2$, so $PRIMES \in RP$

Branching Programs

Take Boolean decision tree, merge equivalent nodes \implies DAG

Def. **Branching Program**

- ▶ DAG, nodes are *query nodes* or two outputs, 0 and 1
- ▶ *query nodes* labeled by variables, 2 outgoing edges (yes/no)
- ▶ designated root (start) node (for evaluation)

Branching programs define the class L
(can build poly-size branching program deciding any language in L)

read-once branching program: on each path from root to 0/1, a variable is queried at most once (not redundant)

$EQ_{ROBP} = \{ \langle B_1, B_2 \rangle \mid B_1 \text{ and } B_2 \text{ are equivalent read-once branching programs} \}$

EQ_{ROBP} is in BPP

Can't randomly choose *boolean* vectors, since programs might differ only on one vector of 2^m

Instead, compute polynomial, starting with 1 at root:

- ▶ YES branch on x_i : multiply with x_i
- ▶ NO branch on x_i : multiply with $(1 - x_i)$

For boolean vector inputs, value is always 0 or 1.

Take product going down on each branch.

Sum all branches entering 1 as the result polynomial.

Equivalent programs \implies equivalent polynomials

If polynomials non-equivalent, probability of difference zero is at most md/f , for m variables, each of degree $\leq d$, over field of size f .
 \implies Since degree of each variable is 1, choose $f > 3m$.