Lecture 10: Algebraic Techniques Fingerprinting, Verifying Polynomial Identities, Parallel Algorithms for Matching Problems

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Overview

- Introduction
 - Why Algebraic Techniques in computer science?
 - Fingerprinting: String equality verification
- Main Problems
 - Polynomial Identity Testing
 - Randomized Matching Algorithms
 - Isolation Lemma
- Remarks
- Acknowledgements

It is hard to overstate the importance of algebraic techniques in computing.

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- many more...

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Suppose Alice and Bob each maintain the same large database of information.¹ They would like to check if their databases are *consistent*.

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Communication complexity setting, randomized algorithms, need to work with high probability.

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Fingerprinting mechanism:

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- what happens when they are different?



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- Choosing p among the first $tn \log(tn)$ primes we have that

$$\Pr[F_p(a) \not\equiv F_p(b)] \le \frac{n}{t n \log t n / \log(t n \log t n)} = \tilde{O}(1/t)$$



Verifying string equality

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• Number of bits sent is $O(\log t + \log n)$. Choosing t = n solves it.

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- **Input:** "Given" two polynomials P(x), Q(x), are they equal?
- \bullet Two polynomials are equal \Leftrightarrow all their coefficients are equal

In string equality, we had

$$P_{A}(x) = \sum_{i=1}^{n} \alpha_{i} x^{i-1} \qquad P_{B} = \sum_{i=1}^{n} b_{i} x^{i-1}$$

where α_{i} , b_{i} $\in \{0,1\}$ (i. $P_{A}(z) \neq P_{B}(z)$ iff $a \neq b$)

wanted $P \in \mathbb{N}$ prime $s + P_{A}(z) \neq P_{B}(z)$ med $P_{A}(z) \neq P_{B}(z)$ and $P_{A}(z) \neq P_{B}(z)$ with more complicated polynomials we may not know whether $P_{A}(z) \neq P_{B}(z)$ for some value of t .

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Technique for string equality testing can be generalized to following setting:

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"Proof: "F[x] is Euclidean domain (so is F[x])

(i.e. "there is division with remainder algorithm")

then induction on degree.

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• Can amplify probability by running multiple times or by choosing larger set *S*.



Lemma (Ore-Schwartz-Zippel-de Millo-Lipton lemma)

Let \mathbb{F} be a field and $P(x_1, \dots, x_n) \in \mathbb{F}[x_1, \dots, x_n]$ be a nonzero polynomial of degree $\leq d$. Then for any set $S \subseteq \overline{\mathbb{F}}$, we have:

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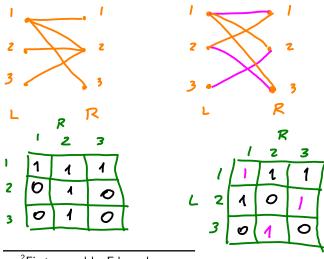
Proof by induction in number of variables.

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• G has perfect matching $\Leftrightarrow \det(X)$ is a non-zero polynomial!²

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- Let $X \in \mathbb{F}^{n \times n}$ be such that

$$X_{i,j} =$$

$$\begin{cases} y_{i,j}, & \text{if there is edge between } (i,j) \in L \times R \\ 0, & \text{otherwise} \end{cases}$$

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$$\det(X) = \sum_{\sigma \in S_n} (-1)^{\sigma} \prod_{i=1}^n X_{i\sigma(i)}$$

- G has perfect matching $\Leftrightarrow \det(X)$ is a non-zero polynomial!²
- Testing if G has a perfect matching is a *special case* of *Polynomial Identity Testing*!

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- Algorithm: evaluate det(X) at a random value for the variables $y_{i,j}$.

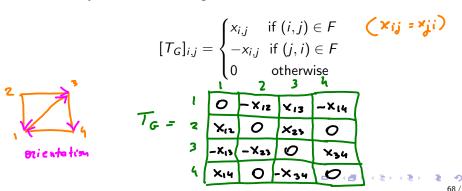


²First proved by Edmonds.

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• Each permutation $\sigma \in S_n$ that yields non-zero term corresponds to a (directed) subgraph of G $H_{\sigma}(V, F_{\sigma})$, where $F_{\sigma} = \{(i, \sigma(i))\}_{i=1}^n$.



$$\sigma = (1234) \rightarrow F_{\sigma} = \{(1,2), (2,3), (3,4), (4,1)\}$$

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- Each vertex in H_{σ} has $|\delta^{out}(i)| = |\delta^{in}(i)| = 1$.

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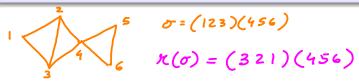
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- If σ only has even cycles, then H_{σ} gives us a perfect matching (by taking every other edge of the graph H_{σ} , ignoring orientation)

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$$\pi(\pi(\sigma)) =$$

• Otherwise, for each $\sigma \in S_n$ (that has <u>odd cycle</u>), there is another permutation $r(\sigma) \in S_n$ that is obtained by reversing odd cycle of H_{σ} containing vertex with *minimum index*.

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- Is there a term that does not cancel? (have to show that $det(T_G) \not\equiv 0$)
- If T_G has a matching, say, $\{1,2\}, \{3,4\}, \ldots, \{2n-1,2n\}$, then take permutation $\sigma = (1\ 2)(3\ 4)\cdots(2n-1\ 2n)$

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- In lecture 21, we will see that we can
 compute the determinant efficiently in parallel

- Introduction
 - Why Algebraic Techniques in computer science?
 - Fingerprinting: String equality verification
- Main Problems
 - Polynomial Identity Testing
 - Randomized Matching Algorithms
 - Isolation I emma
- Remarks
- Acknowledgements

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Remark

The isolation lemma could be quite counter-intuitive. A set system can have $\Omega(2^n)$ sets. On average, there are $\Omega(2^n/(2n^2))$ sets of a given weight, as max weight is $\leq 2n^2$. Isolation lemma tells us that with high probability there is *only one* set of minimum weight.

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- α_v is *independent* of w(v), and w(v) chosen uniformly at random from [2n].
- **③** $Pr[v \text{ ambiguous}] \le 1/2n \Rightarrow_{\text{union bound}} Pr[∃ \text{ ambiguous element}] \le 1/2$



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 - lacktriangle Probability that this happens is $\leq 1/2$. (step 8)



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- many more...

Derandomizing (i.e., obtaining deterministic algorithms) for some of these settings (whenever possible) is *major open problem* in computer science.

Potential Final Projects

- Can we derandomize the perfect matching algorithms from class?
- A lot of progress has been made in the past couple years on this question in the works [Fenner, Gurjar & Thierauf 2019] and subsequently [Svensson & Tarnawski 2017]
- Survey of the above, or understanding these papers is a great final project!

Acknowledgement

- Lecture based largely on:
 - Lap Chi's notes
 - [Motwani & Raghavan 2007, Chapter 7]
 - [Korte & Vygen 2012, Chapter 10].
- See Lap Chi's notes at https://cs.uwaterloo.ca/~lapchi/cs466/notes/L07.pdf

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SIAM Journal on Computing



Svensson, Ola and Jakub Tarnawski (2017)

The matching problem in general graphs is in quasi-NC.

IEEE 58th Annual Symposium on Foundations of Computer Science