## Bipartite Perfect Matching is in Quasi-NC

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January 19, 2025

## Outline

### **Abstract**

This is a summary of "Bipartite Perfect Matching is in Quasi-NC" by Fenner, Gurjar, and Thierauf.

- Introduction
- 2 Combinatorics and Probability
- Isolation
- 4 Cycles and Circulation
- The Theorem

## **Notation**

#### Notation

Throughout this presentation, *ALL* the graphs are:

- undirected
- bipartite
- balanced
- labeled

Usually we call our graph G and the set of edges E.

The set of edges can be regarded as a relation  $E \subseteq [n] \times [n]$ .

#### Definition

- We define  $\mathcal{NC}^k$  to be the class of problems that can be solved by a polynomial-time uniform family of circuits of polynomial size and depth  $\mathcal{O}(\log^k n)$ .
- Equivalently,  $\mathcal{NC}^k$  is the class of problems that can be solved by a polynomial number of processors in  $\mathcal{O}(\log^k n)$  time.

•

$$\mathcal{NC} = \bigcup_{k \geq 0} \mathcal{NC}^k$$

#### **Fact**

DET, the problem of computing the determinant of an  $n \times n$ matrix with poly(n)-bounded entries, is in  $\mathcal{NC}^2$ .

## Perfect Matching

#### Definition

- A *perfect matching* of a graph *G* is a set of edges such that every vertex is incident to exactly one edge.
- The perfect matching problem is to determine whether a graph has a perfect matching, or to find one.
- The decision version of the perfect matching problem is denoted PM.
- The search version of the perfect matching problem is denoted SearchPM.

Introduction

It has already been known that perfect matching (whether decision or search) can be solved in randomized  $\mathcal{NC}$ .

## Open Problem

Can perfect matching be solved in  $\mathcal{NC}$ ?

## Today

There exist algorithms in:

- Quasi $-\mathcal{NC}^2$  for PM (that is, with  $\mathcal{O}(n^{\log n})$  processors and  $\mathcal{O}(\log^2 n)$  depth)
- randomized  $\mathcal{NC}^2$  for PM with only  $\mathcal{O}(\log^2 n)$  random bits.
- (We will not see this) Quasi $-\mathcal{NC}^2$  for PM (that is, with  $\mathcal{O}(n^{\log n})$  processors and  $\mathcal{O}(\log^2 n)$  depth)
- (We will not see this)  $\mathcal{NC}^3$  for SearchPM with only  $\mathcal{O}(\log^2 n)$ random bits.

- The bi-adjacency matrix:  $\mathbf{A}_G = \mathbf{A}_E = A$  of G (or E) is an  $n \times n$  matrix where  $A_{ij} = 1_{(i,j) \in E}$
- We write  $B \leq C$  iff  $\forall i, j.B_{ij} \leq C_{ij}$
- A permutation matrix is a matrix  $\mathbf{A}_{\Gamma}$  where  $\Gamma = \{(i, \sigma i) : i \in [n]\}$  for some permutation  $\sigma \in \mathcal{S}_n$

## Exercise

For which  $E \subseteq [n] \times [n]$  is  $\mathbf{A}_E$  a permutation matrix?

#### Exercise

For which  $B \in \mathbb{R}_{>0}^{n \times n}$  and E does  $B \leq \mathbf{A}_E$  hold?

- A weight function is a function  $w: E \to \mathbb{N}$ .
- We extend it naturally to a function  $w: 2^E \to \mathbb{N}$  by setting  $w(S) = \sum_{e \in S} w(e)$  for all  $S \subseteq E$ .
- Even more generally (why?), we extend it to a *linear function*  $w: \mathbb{R}^{n \times n} \to \mathbb{R}$  by setting  $w(A) = \sum_{ij \in E} w(ij)A_{ij}$  for all  $A \in \mathbb{R}^{n \times n}$ .

## Definitions (Perfect Matching)

- A *perfect matching* is a set of edges *M* such that every vertex is incident to exactly one edge in *M*.
- Equivalently, M is a perfect matching iff  $\mathbf{A}_M$  is a permutation matrix.

# • A matrix $M \in \mathbb{R}_{>0}^{n \times n}$ is doubly stochastic if the sum of the

- entries in each row and column is 1.
- Equivalently, M is doubly stochastic if  $M\mathbf{1} = M^T\mathbf{1} = \mathbf{1}$ .

## Definition (Convex Hull)

The convex hull of a set S is the smallest convex set containing S, denoted CH(S).

It is precisely the set of all convex combinations of elements of S,

i.e., 
$$\mathbf{CH}(S) = \left\{ \sum_{i=1}^{k} \lambda_i x_i : x_i \in S, \lambda_i \geq 0, \sum_{i=1}^{k} \lambda_i = 1 \right\}.$$

## Birkhoff's Theorem II

## Theorem (Birkhoff, 1946)

The set of doubly stochastic matrices, denoted  $\mathcal{B}_n$ , is the convex hull of the permutation matrices.

### Hint

Use Hall's marriage theorem.

## Perfect Matching Polytope

## Theorem (Birkhoff, 1946)

The set of doubly stochastic matrices is the convex hull of the permutation matrices.

## Definition (Perfect Matching Polytope)

The perfect matching polytope  $\mathcal{P}_G$  is the convex hull of the bi-adjacency matrices of all the perfect matchings of G.

## Corollary

The perfect matching polytope  $P_G$  is exactly the matrices B in the Birkhoff polytope  $B_n$  such that  $B \leq \mathbf{A}_G$ :

$$\mathcal{P}_G = \{B : B \in \mathcal{B}_n \land B \leq \mathbf{A}_G\}$$

#### Definition

Let M be a perfect matching. We define its  $sign \operatorname{sgn} M = \operatorname{sgn} \sigma$  where  $\sigma$  is the permutation given by M.

#### Definition

Let  $\{w_i : E \to \mathbb{N}\}_{i \in [k]}$  be weight functions.

We define the weighted Edmonds matrix  $D^{E,w}$  as follows:

$$\left(D^{E,w}\right)_{ij} = \begin{cases} x_{ij}^{(k)^{w_k(ij)}} & \text{if } (i,j) \in E, \\ 0 & \text{otherwise.} \end{cases}$$

Let  $a \in \mathbb{N}^k$  be a vector of variables. We also denote  $D_a^{E,w} = D^{E,w}(x_{ii}^{(k)} := a_k)$ 

#### **Definitions**

- A weight function  $w: E \to \mathbb{N}$  is called *isolating* for E if the existence of a perfect matching in E implies that E has a unique minimum weight perfect matching, which is then called *isolated*.
- Similarly, a sequence of weight functions  $w = (w_i)_{i \in [k]}$  is called isolating for E if the existence of a perfect matching in E implies that E has a unique minimum weight perfect matching with respect to  $w = (w_0, \ldots, w_k)$ , ordered lexicographically.

## Corollary

- If E has no perfect matchings, then  $\det D^{E,w} = 0$ .
- If w is isolating for E, then  $\det D^{E,w} \neq 0 \iff$  there is a perfect matching.



Introduction

Let  $E = \{e_k\}_k$  be an enumeration of the edges. Let  $w : E \to \mathbb{N}$  be defined by  $w(e_k) = 2^k$ . Every subset of E has a unique weight. In particular, w is isolating for E. Is this enough?

## Isolation Lemma

## Lemma (Mulmuley, Vazirani & Vazirani 1987)

Let B be a finite set.

Let k be a positive integer.

Let  $\mathcal{F} \subseteq 2^B$  be a nonempty family of subsets of B.

Let  $w : B \to k$  be a be chosen uniformly at random among all functions from B to [k].

Then, with probability at least 1 - |B|/k, there exists a unique set  $S \in \mathcal{F}$  with a minimum weight among all sets in  $\mathcal{F}$ .

## Corollary

Let  $w: E \to [n^3]$  be a random weight function.

Then with probability at least 1-1/n, w is isolating for G.

#### Proof.

Take  $\mathcal{F}$  as the family of perfect matchings of G.

As,  $|E| \le n^2$ , by the isolation lemma, with probability at least

$$(1-|E|/n^3) \ge 1-n^2/n^3 = 1-1/n$$

w is isolating for G.

## Corollary

There exists an RNC algorithm that can solve PM, which uses poly(n) random bits.



## Circulation

## Definition

Let  $w: E \to \mathbb{N}$  be a weight function. The *w-circulation* of a cycle  $C = e_1, \ldots, e_{2k}$  is defined as

$$c_w(C) = |w(e_1) - w(e_2) + \ldots + w(e_{2k-1}) - w(e_{2k})|$$

This is well defined because we take the absolute value.

### Remark

Let  $M_1$  and  $M_2$  be two perfect matchings of G, and suppose  $C \subseteq M_1 \triangle M_2$ .

Then 
$$c_w(C) = |w(C \cap M_1) - w(C \cap M_2)|$$
.

### Claim

The symmetric difference of two perfect matchings is a union of disjoint cycles.

Each cycle consists of interleaved edges from the two matchings.

#### Proof.

The degree of each vertex is 1 in a perfect matching.

Thus, in the symmetric difference of two perfect matchings, each vertex has degree 0 or 2.

Therefore it is a union of disjoint cycles. The rest is easy to see.



#### Lemma

Let  $w: E \to \mathbb{N}$  be a weight function.

If every cycle in G has a nonzero circulation, then w is isolating for G.

#### Proof.

On the contrary, suppose that  $M_1$  and  $M_2$  are two minimum perfect matchings of G. Choose some cycle  $C \subseteq M_1 \triangle M_2$ . Since  $c_w(C) > 0$ , we have  $w(C \cap M_1) \neq w(C \cap M_2)$ . WLOG, assume that  $w(C \cap M_1) < w(C \cap M_2)$ . Then  $M_2 \triangle C$  is a perfect matching with weight less than  $w(M_2) = w(M_1)$ , contradicting minimality.

#### Definition

Suppose G has a perfect matching. Let  $w: E \to \mathbb{N}$  be a weight function.

Then  $G_w \subseteq G$  is the union of all minimum weight perfect matchings of G, relatively to w.

#### Theorem

Let  $G^0 = G$  be and  $G^{i+1} = (G^i)_{w_i}$ .

Let  $k \ge \log n - 2$  and let  $w_0, \dots, w_k : E \to \mathbb{N}$  be weight functions such that  $w_i$  gives nonzero circulation to all the cycles of size at most  $4 \cdot 2^i$  in  $G_i$ .

Then the joint weight function  $w = (w_0, \dots, w_k)$  is isolating.

## Lemma

Let  $r \ge 2$  be even. Suppose G has no cycles of size at most 2r. Then G has at most  $n^4$  cycles of size at most 4r.

#### Proof.

Introduction

- Equivalently, there is at most one path of length  $\leq r$  between any two vertices.
- Let  $v_0, v_1, v_2, v_3$  be vertices in  $V_1$ , such that the distance between  $v_i$  and  $v_{i+1 \mod 4}$  is  $\leq r$ .
- There exists a partial function from  $V_1^4$  to cycles in G that maps  $(v_0, v_1, v_2, v_3)$  to the cycle composed of the unique paths of length  $\leq r$  between  $v_i$  and  $v_{i+1 \mod 4}$ .
- Let C be a cycle of size ≤ 4r.
   We can choose arbitrarily v<sub>0</sub>, v<sub>1</sub>, v<sub>2</sub>, v<sub>3</sub> ∈ C such that the distance between v<sub>i</sub> and v<sub>i+1 mod 4</sub> is ≤ r.
- This function is onto the set of cycles of size  $\leq 4r$ . Thus, there are at most  $|V_1|^4 = n^4$  cycles of size  $\leq 4r$ .



#### Lemma

Suppose G has a perfect matching and let  $w : E \to \mathbb{N}$  be a weight function.

Then every perfect matching in  $G_w$  has the same weight as every minimum weight matching in G.

#### Proof.

If w is isolating,  $G_w$  is just a single perfect matching.

Otherwise, suppose the minimum weight for perfect matchings in G is m. Suppose there exist t such matchings and let X be their average. Then, w(X) = m.

Let N be a perfect matching in  $G_w$ . Every edge is contained in some minimum weight perfect matching, so  $X-\frac{1}{t}N\geq 0$ . It can easily be seen that the sum of every row and column is (t-1)/t and thus the matrix  $\frac{t}{t-1}X-\frac{1}{t-1}N$  is doubly stochastic. Therefore, it lies in  $\mathcal{P}_G$  and thus has weight at least m. That is.

$$m \le w\left(\frac{t}{t-1}X - \frac{1}{t-1}N\right) = m \cdot \frac{t}{t-1} - w(N) \cdot \frac{1}{t-1}$$

From that it follows that  $w(N) \leq m$ .



#### Lemma

Suppose G has a perfect matching and let  $w: E \to \mathbb{N}$  be a weight function.

Then the w-circulation of every cycle in  $G_w$  is zero.



#### Proof.

Introduction

Let X be the average of all perfect matchings in  $G_w$ . Let t be the number of perfect matchings,  $\varepsilon = 1/t$ , and let  $M_1, \ldots, M_t$  be the perfect matchings. Then  $X = \varepsilon \sum_{i=1}^t M_i$ . Since each edge is contained in some perfect matching, we have  $X_{ii} \geq \varepsilon$  for all i, j such that  $(i, j) \in E$ . Let  $C = e_1, \ldots, e_{2p}$  be a cycle in H. Define Y by

$$Y_{ij} = \begin{cases} X_{ij} + (-1)^k \varepsilon & \text{if } (i,j) = e_k \in C, \\ X_{ij} & \text{otherwise.} \end{cases}$$

Then clearly Y > 0 and  $Y\mathbf{1} = Y^T\mathbf{1} = \mathbf{1}$ . Therefore, Y lies in the perfect matching polytope.

Since all the perfect matchings have the same weight, w(Y) = w(X) and thus w(Y - X) = 0. But  $c_w(C) = \varepsilon w(Y - X)$ , and thus  $c_w(C) = 0$ .

#### Lemma

Using  $\mathcal{O}(\log n)$  random bits we can generate a weight assignment  $w: E \to \mathbb{N}$  with  $w \leq \operatorname{poly}(n)$  such that for every set of  $n^4$  cycles, w gives nonzero circulation to all of them with probability at least 1-1/n.

#### Proof.

Let  $w(e_k) = 2^k$ . Let the cycles be  $C_1, \ldots, C_s$ . Then,  $c_w(C_i) \neq 0$  for all i is equivalent to  $\prod_{i=1}^{n^4} c_w(C_i) \neq 0$ . This product is bounded by  $poly(n)^{n^4} = 2^{poly(n)}$ .

Thus, it has at most poly(n) prime factors, say k prime factors. Choose t = kn.

Then if we choose a random prime amongst the first t primes  $[p_1, \ldots, p_t]$ , then if the product is nonzero, with probability at least 1 - 1/n it is still true modulo the chosen prime.



### Theorem

Let  $k \ge \log n - 2$  and let  $w_0, \ldots, w_k : E \to \mathbb{N}$  be weight functions. Let  $G^0 = G$  be and  $G^{i+1} = (G^i)_{w_i}$ .

Suppose that for every set of  $w_i$  gives nonzero circulation to all the cycles of size at most  $4 \cdot 2^i$  in  $G_i$ .

Then the joint weight function  $w = (w_0, ..., w_k)$  is isolating.

### Proof.

Since  $w_i$  gives nonzero circulation to all the cycles in  $G^i$ , all those cycles do not appear in  $G^{i+1}$ .

In particular,  $G^k$  has no cycles of size at most  $4 \cdot 2^k \ge n$ .

That is,  $G^k$  has a unique perfect matching. We want to show that it is isolated by w. This follows from the next lemma.

## Lemma

Suppose  $M_1$  is a matching that appears in  $G^i$  but not in  $G^{i+1}$  and  $M_2$  is a matching that appears in  $G_{i+1}$ . Then  $w_i(M_2) < w_i(M_1)$ .

## Proof.

Since  $M_2$  appears in  $G_{i+1}$ , it has the same weight as any  $w_i$ -minimum weight matching in  $G_i$ . That is  $w_i(M_2) \leq w_i(M_1)$ . But if  $w_i(M_1) = w_i(M_2)$  held, then  $M_1$  would have been in  $G^{i+1}$ by its definition.



## Deciding PM in randomized $\mathcal{NC}^2$

The following algorithm solves PM in randomized  $\mathcal{NC}^2$  with  $\mathcal{O}(\log^2 n)$  random bits.

## **Algorithm 1** Decide PM (randomized)

```
Let k = \log n - 2.
```

Generate random weight functions  $w_0, \ldots, w_k$ .

Generate random numbers  $r_0, \ldots, r_k$ .

Create  $w_0, \ldots, w_k$ .

Combine  $w = (w_0, \ldots, w_k)$ .

Calculate det  $D_r^{G,w}$ .

**return** det  $D_r^{G,w} \neq 0$ 



## Deciding PM in randomized Quasi- $\mathcal{NC}^2$

The following algorithm solves PM in Quasi- $\mathcal{NC}^2$ .

## **Algorithm 2** Decide PM (deterministic)

```
Let k = \log n - 2. for all Possible weight functions w_0, \ldots, w_k and r, simultaneously do

Calculate \det D_r^{G,w}.

if \det D_r^{G,w} \neq 0 then

return True

end if

return False

end for
```

## Questions?

