Matching Size and Matrix Rank Estimation in Data Streams

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Joint work with Sanjeev Khanna (Penn), and Yang Li (Penn)

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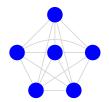
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- Input is presented as a data stream, for instance, as a sequence of edges in case of a graph input.
- Algorithm sees the entire input once and only has a small space to store information about the input as it passes by.
- At the end of the sequence, the algorithm outputs a solution using only the stored information.

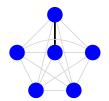
A graph stream:

Stream:



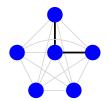
A graph stream:

Stream: $+e_1$



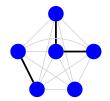
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Stream: $+e_1, +e_7$



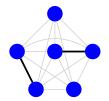
A graph stream:

Stream: $+e_1, +e_7, +e_{11}$



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Stream: $+e_1, +e_7, +e_{11}, -e_1$



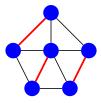
Two relevant models for our purpose:

• Insertion-Only Streams. Only contains positive updates.

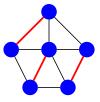
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- Insertion-Only Streams. Only contains positive updates.
- Dynamic Streams. Contains both positive and negative updates.

• Matching: A collection of vertex-disjoint edges.

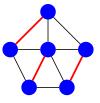


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• Perfect Matching: Every vertex is in the matching.

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Maximum Matching problem: Find a matching with a largest number of edges.

Maximum matching is a fundamental problem with many applications.

- Many celebrated algorithms in the classical setting: Ford-Fulkerson, Edmond's, Hopcroft-Karp, Mucha-Sankowski, Madry's, . . .
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This talk: sublinear space algorithms for the matching problem in the streaming model.

Finding a Matching vs Estimating Size

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Are there any qualitative difference in the space needed to achieve these goals?

Arguably the most studied problem in the graph streaming literature.

[McGregor, 2005] [Feigenbaum et al., 2005] [Eggert et al., 2009] [Epstein et al., 2011] [Goel et al., 2012] [Konrad et al., 2012] [Zelke, 2012] [Ahn et al., 2012] [Ahn and Guha, 2013] [Guruswami and Onak, 2013] [Kapralov, 2013] [Kapralov et al., 2014] [Crouch and Stubbs, 2014] [McGregor, 2014] [Chitnis et al., 2015] [Ahn and Guha, 2015] [Esfandiari et al., 2015] [Konrad, 2015] [Bury and Schwiegelshohn, 2015] [Assadi et al., 2016] [Chitnis et al., 2016] [McGregor and Vorotnikova, 2016b] [Esfandiari et al., 2016] [Paz and Schwartzman, 2017] [Ghaffari, 2017] [Kale et al., 2017] ...

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Dynamic streams:

- $\Omega(n^2/\alpha^3)$ space is necessary for α -approximation [Assadi et al., 2016].

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 - Random arrival insertion-only streams [Kapralov et al., 2014].
 - ► Bounded arboricity graphs [Esfandiari et al., 2015] ...
- Lower bounds (Insertion-only streams):
 - (1 + ε)-approximation requires Ω(n^{1−O(ε)})
 space [Esfandiari et al., 2015, Bury and Schwiegelshohn, 2015].
 - Deterministic α-approximation requires Ω(n/α) space [Chakrabarti and Kale, 2016].

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Theorem

There is a randomized algorithm that outputs an α -approximate estimate of maximum matching size in:

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In constrast, to find an α -approximate matching, the space necessary is:

- $\Omega(n/\alpha)$ in insertion-only streams.
- $\Omega(n^2/lpha^3)$ in dynamic streams.

Algorithms for α -Estimation of Matching Size

The main ingredient of our algorithms is the following sampling lemma:

Lemma (Vertex Sampling Lemma)

Let *H* be a subgraph of *G* obtained by sampling each vertex independently w.p. $1/\alpha$. Define:

 μ_G : the maximum matching size in G,

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 $\frac{\mu_G}{\alpha^2} \le \mu_H \le \frac{2\mu_G}{\alpha}.$

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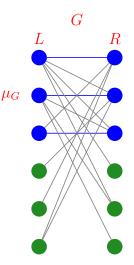
$$\frac{\mu_G}{\alpha^2} \le \mu_H \le \frac{2\mu_G}{\alpha}.$$

Therefore, maximum matching size in H is an α -estimation for the maximum matching size in G.

Proof by Picture

Any graph G with a maximum matching size of μ_G looks as follows:

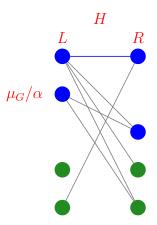
- A matching of size μ_G between the blue vertices.
- No edges between the green vertices.



Proof by Picture

The vertex sampled graph H then look as follows:

- A matching of size μ_G/α^2 between the blue vertices $\implies \mu_H \ge \mu_G/\alpha^2$.
- All edges are incident on μ_G/α blue vertices $\implies \mu_H \le 2\mu_G/\alpha.$



An α -Estimation Algorithm

To distinguish between graphs with maximum matching of size $\geq k$ and $o(k/\alpha)$:

- **1** Sample each vertex in G w.p. $1/\alpha$ to obtain H.
- 2 Test whether H has a matching of size at least $\Omega(k/\alpha^2)$ or not.

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Can be implemented in:

- $\tilde{O}(k/\alpha^2) = \tilde{O}(n/\alpha^2)$ in insertion-only streams.
- $\tilde{O}(k^2/\alpha^4) = \tilde{O}(n^2/\alpha^4)$ in dynamic streams.

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We make progress on each of these questions.

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RS(n) denotes the maximum number of edge-disjoint induced matchings of size $\Theta(n)$ in an *n*-vertex graph:

[Fischer et al., 2002] $n^{\Omega(1/\log \log n)} \leq \mathsf{RS}(n) \leq n/\log n$ [Fox et al., 2015]

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Any randomized α -approximate estimate of maximum matching size requires $\Omega(n/\alpha^2)$ in dynamic streams. Furthermore, even if we restrict to sparse graphs with arboricity $O(\alpha)$, $\Omega(\sqrt{n}/\alpha^{2.5})$ space is necessary.

There is an active line of research on estimating matching size of bounded arboricity graphs in graph streams [Chitnis et al., 2016] [Bury and Schwiegelshohn, 2015] [Esfandiari et al., 2015] [McGregor and Vorotnikova, 2016b] [Cormode et al., 2016] [McGregor and Vorotnikova, 2016a] ...

Schatten *p*-Norms

Given an $n \times n$ matrix A, for any $p \in [0, \infty)$:

Schatten *p*-norm of *A* is the *p*-th frequency moment of vector of singular values $(\sigma_1, \ldots, \sigma_n)$ of *A*.

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- $||A||_0 = \mathsf{Rank} \text{ of } A.$
- $||A||_1 = \text{Trace norm of } A.$
- $||A||_2 =$ Frobenius norm of A.
- $||A||_{\infty} = \text{Operator norm of } A.$

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Previous work:

- For p = 0, Ω(n^{1-g(ε)}) space is necessary [Bury and Schwiegelshohn, 2015].
- For p ∈ (0,∞) \ 2ℤ, Ω(n^{1-g(ε)}) space is necessary [Li and Woodruff, 2016].
- For p ∈ 2Z \ {0}, Ω(n^{1-2/p}) space is necessary [Li and Woodruff, 2016] (and is sufficient for sparse matrices).

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We answer this question for the case of rank computation.

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It is well-known that computing maximum matching size of a graph is equivalent to computing the rank of the (symbolic) Tutte matrix.

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As a corollary, all our lower bounds for matching size estimation also extend to the matrix rank computation problem. In particular,

- An $\Omega(n^{2-O(\varepsilon)})$ space lower bound for $(1+\varepsilon)$ -estimation of rank in dense matrices.
- An $\tilde{\Omega}(\sqrt{n})$ space lower bound for any $\operatorname{polylog}(n)$ -estimation of rank in sparse matrices.

An $\Omega(n^{2-O(\varepsilon)})$ Lower Bound for Dynamic Streams

Theorem

Any randomized $(1 + \varepsilon)$ -approximate estimation of maximum matching size requires $\Omega(n^{2-O(\varepsilon)})$ space in dynamic streams.

Consider the following two-player one-way communication problem.

MAXMATCHING:

- Alice is given a matching M on vertices V.
- **2** Bob is given a collection of edges E_B on vertices V.
- Ice sends a single message to Bob and Bob outputs an estimation of maximum matching size in $G(V, M \cup E_B)$.

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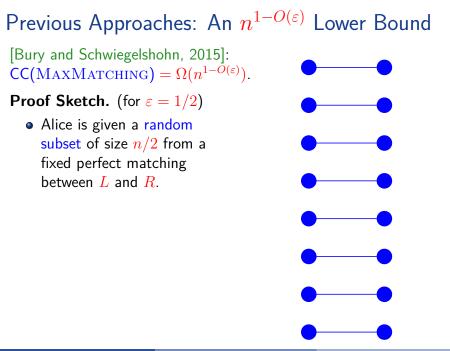
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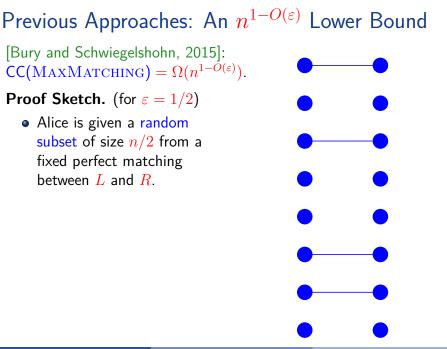
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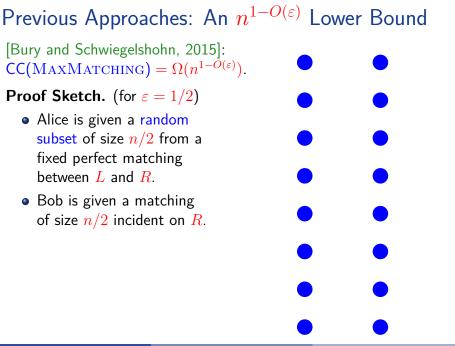
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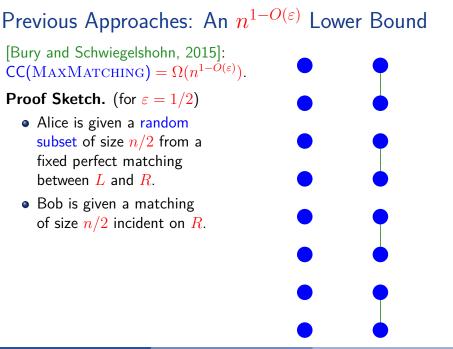
Fact. $CC(MAXMATCHING) \leq$ space complexity of any streaming algorithm for estimating maximum matching size.

[Bury and Schwiegelshohn, 2015]: $CC(MAXMATCHING) = \Omega(n^{1-O(\varepsilon)}).$









Previous Approaches: An $n^{1-O(\varepsilon)}$ Lower Bound [Bury and Schwiegelshohn, 2015]: $\mathsf{CC}(\mathsf{MAXMATCHING}) = \Omega(n^{1-O(\varepsilon)}).$ **Proof Sketch.** (for $\varepsilon = 1/2$) • Alice is given a random subset of size n/2 from a fixed perfect matching between L and R_{\star} Bob is given a matching of size n/2 incident on R. • Yes case: Each edge of Bob's matching is incident on even number of Alice's matching \implies MAXMATCHING = 3n/4.

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problem of [Gavinsky et al., 2007].

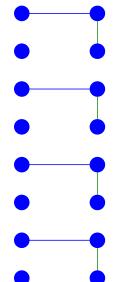
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Proof Sketch. (for $\varepsilon = 1/2$)

- A better than 3/2-approximation distinguishes between the two cases.
- Distinguishing between the two cases requires Ω(√n) communication by a reduction from the boolean hidden matching problem of [Gavinsky et al., 2007].





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The hope is that communication complexity of this problem is now $\geq t \cdot CC(MAXMATCHING)$.

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Formalizing the lower bound \implies a direct-sum style argument using information complexity.

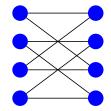
Sepehr Assadi (Penn)

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A graph G(V, E) whose edges can be partitioned into t induced matchings of size r each.

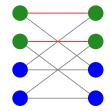
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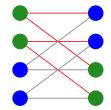
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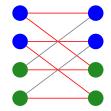
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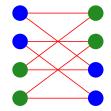
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Theorem ([Fischer et al., 2002])

There exists an (r, t)-RS graph on n vertices with $t = n^{\Omega(1/\log \log n)}$ induced matchings of size $r = (1 - \varepsilon) \cdot n/4$.

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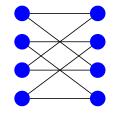
Theorem ([Fischer et al., 2002])

There exists an (r, t)-RS graph on n vertices with $t = n^{\Omega(1/\log \log n)}$ induced matchings of size $r = (1 - \varepsilon) \cdot n/4$.

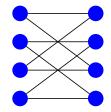
Theorem ([Alon et al., 2012])

There exists an (r, t)-RS graph on n vertices with $t = n^{1+o(1)}$ induced matchings of size $r = n^{1-o(1)}$.

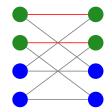
• Let G_1 be an (r, t)-RS bipartite graph on n vertices on each side.



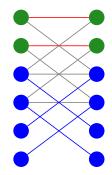
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- To Alice, we give random subset of size r/2 from each induced matchings M_1, \ldots, M_t of G_1 .



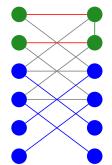
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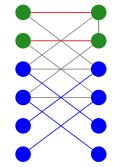


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 - A graph E_B over the set of vertices in M_j★.



• Size of the maximum matching in this graph:

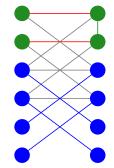
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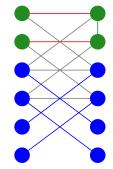
• For $r = \Theta(n)$, Alice and Bob need to solve MAXMATCHING (M_{j^*}, E_B) for $(1 + \varepsilon)$ -approximation.



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- To solve this for an unknown matching M_{j^*} , the message length must be $\geq t \cdot CC(MAXMATCHING)$.



Main limitation of this approach:

- Requires $r = \Theta(n) \implies t = \mathsf{RS}(n)$.
- $\mathsf{RS}(n)$ maybe as large as $n/\log n$.
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We bypass the $r = \Theta(n)$ limitation in dynamic streams using the characterization result of [Li et al., 2014, Ai et al., 2016] in terms of the simultaneous communication complexity.

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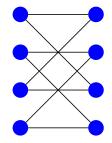
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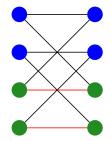
[Ai et al., 2016]: Communication lower bounds in this model imply identical space lower bound for dynamic streaming algorithms.

• Each player is given an (r, t)-RS graph on N vertices.



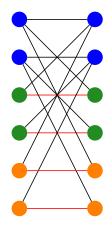
Local view of P^i

- Each player is given an (r, t)-RS graph on N vertices.
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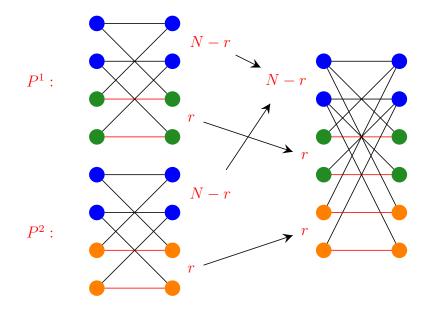


$\begin{array}{c} \text{Special matching} \\ \text{of } P^i \end{array}$

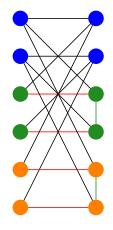
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Global view



- Each player is given an (r, t)-RS graph on *N* vertices.
- One of the induced matching $M_{j^{\star}}^{(i)}$ of each player $P^{(i)}$'s graph is special, unknown to the player.
- Across the players, vertices in the special matchings are unique, while other vertices are shared.
- To the referee, we provide k subgraphs $E_B^{(1)}, \ldots, E_B^{(k)}$ such that each pair $(M_{j^\star}^{(i)}, E_B^{(i)})$ forms the same instance of MAXMATCHING.

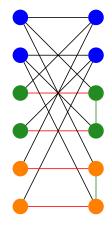


Global view

- Define MAXMATCHING $(M, E_B) :=$ MAXMATCHING $(M_{j^{\star}}^{(1)}, E_B^{(1)}) = \ldots =$ MAXMATCHING $(M_{j^{\star}}^{(k)}, E_B^{(k)}).$
- The maximum matching size in G is:

 $\approx 2(N-r) + k \cdot \text{MaxMatching}(M, E_B)$

• For $r = N^{1-o(1)}$ and $k \approx \frac{1}{\varepsilon} \cdot N^{o(1)}$, MAXMATCHING (M, E_B) is the dominating term for $(1 + \varepsilon)$ -approximation.



Global view

• The players need to solve $MAXMATCHING(M, E_B)$.

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- To conclude,

$$N^{1-O(\varepsilon)} \le k \cdot s/t \implies s \ge \frac{1}{k} \cdot t \cdot N^{1-O(\varepsilon)} \approx N^{2-O(\varepsilon)}$$

as $t = N^{1+o(1)}$ by [Alon et al., 2012] and $k = \Theta_{\varepsilon}(N^{o(1)})$.

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Theorem

Any randomized $(1 + \varepsilon)$ -approximate estimation of maximum matching size requires $\Omega(n^{2-O(\varepsilon)})$ space in dynamic streams.

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 - $\Omega(n/\alpha^2)$ space is necessary vs. $\widetilde{O}(n^2/\alpha^4)$ space is sufficient.
- Similar-in-spirit lower bounds for Schatten p-norms for p > 0?

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