# A Family of Approximation Algorithms for the Maximum Duo-Preservation String Mapping Problem

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July 13, 2017

## Minimum Common String Partition

Input: two strings X, Y, where Y is a permutation of X. Output: partition of X into the least number of pieces that can be rearranged (without reversing) and concatenated to obtain Y.

$$X$$
: xyzabcbxy

$$Y:$$
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$$X: \quad \boxed{x \ y \ z} \boxed{a \ b} \boxed{c} \boxed{b} \boxed{x \ y}$$

$$Y:$$
 a b b c x y z x y

## Hardness of MCSP

Goldstein, Kolman, Zheng ['04]

MCSP is APX-hard.

Cormode, Muthukrishnan ['07]

Almost linear-time  $O(\log n \cdot \log^* n)$ -approximation algorithm.

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# Maximum Duo-Preservation String Mapping Problem

Complementary problem: maximize number of **duos** – consecutive letters not split apart.

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$$Y: \quad |\mathbf{a} \ \mathbf{b}| |\mathbf{c}| |\mathbf{x} \ \mathbf{y} \ \mathbf{z}| |\mathbf{x} \ \mathbf{y}|$$

$$|X| = \#duos + \#pieces$$

Easier? No:

Boria, Kurpisz, Leppänen, Mastrolilli ['14]:

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Preserved duos: xy, yz, ab, xy

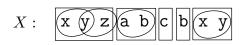
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## Results for MPSM

Authors	year	ratio
Boria et al.	'14	4
Boria et al.	'16	3.5
Brubach	'16	3.25
Xu et al.	'17	2.917
DGO-N	'17	$2+\varepsilon$

Boria, Kurpisz, Leppänen, Mastrolilli ['14]: It is NP-hard to approximate MPSM within 1.00042  $-\varepsilon$  for every  $\varepsilon>0$ .

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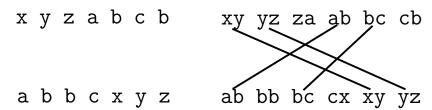
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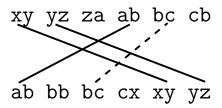
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## Graph representation

Bipartite graph with nodes – duos in both strings:

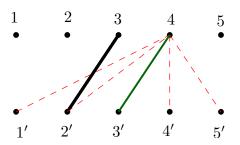


Maximum consecutive bipartite matching:



# Maximum Consecutive Bipartite Matching

When we take the edge (2',3) to the matching:



## **Definitions**

#### Streak:



#### Conflicting edges



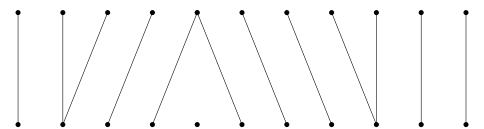
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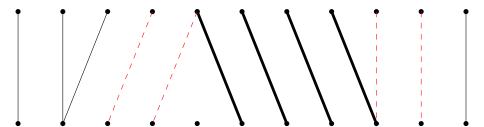
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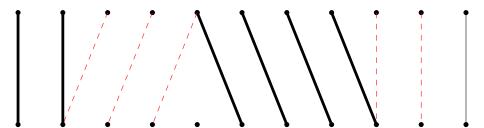


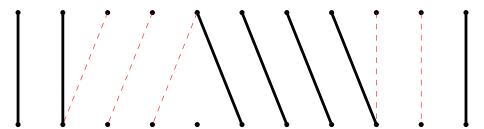
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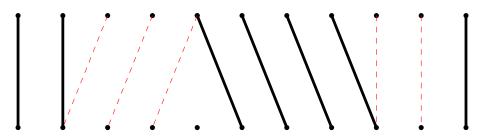








As long as possible take the longest possible streak from G.



If we stop the algorithm when streaks contain less than k edges:

#### Lemma

There are  $(2 + \frac{2}{k}) \cdot |GREEDY|$  edges from optimal solution conflicting with GREEDY.

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#### Lemma

Streaks of length at least k:  $(2 + \frac{2}{k})$ -approximation.

#### Streaks smaller than k need another phase:

- k = 1 The greedy algorithm alone yields 4-approximation.
- k = 2 Maximum matching for the remaining edges yields 3-approximation.
- k = 3 Local search technique used by Boria et al. yields 2.67-approximation.

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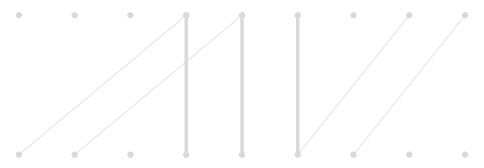
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# $(2 + \varepsilon)$ -approximation

## BOUNDEDSIZEIMPROVEMENTS(t)

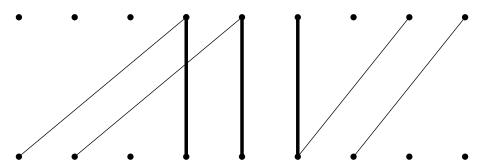
- try every subset  $E_{add}$ ,  $E_{del}$  of at most t edges
- if  $|E_{del}| < |E_{add}|$ , try  $ALG \setminus E_{del} \cup E_{add}$



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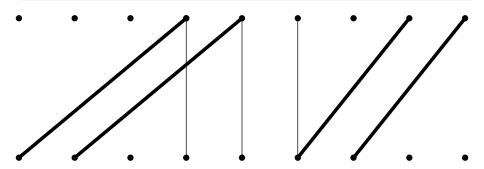
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## Final algorithm

- **1** run Greedy for  $k = \left\lceil \frac{2}{\varepsilon} \right\rceil$
- ② run BoundedSizeImprovements( $\lceil \frac{4}{\varepsilon} \rceil + 1$ ).

#### **Theorem**

Combining the greedy algorithm with local improvements yields a  $(2+\varepsilon)$ -approximation for MCBM in  $n^{O(1/\varepsilon)}$  time, for any  $\varepsilon>0$ .

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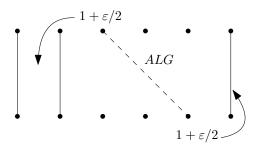
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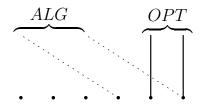
Proof: assign  $2 + \varepsilon$  credits to every edge from *ALG* 



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#### Credit distribution scheme

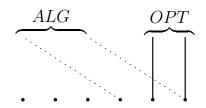
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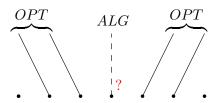
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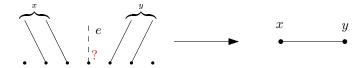
#### Balance of a streak

Balance: number of credits distributed to a streak minus its size.

#### Lemma

Balance of every streak is at least -2.

New graph with nodes – streaks of OPT:



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Every connected component of the new graph has overall balance at least -1.

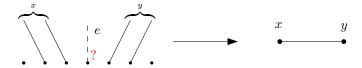
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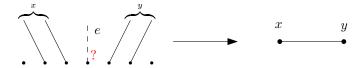
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- enough to cover 1 missing credit.

## Small components

- not greater than t,
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# Open problems

- What are the actual bounds for the problem?
  - ▶ lower bound:  $1.00042 \varepsilon$
  - upper bound:  $2 + \varepsilon$
- Is k-MPSM significantly easier than MPSM?

Xu, Chen, Luo, Lin ['17]:

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