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## Tree-Constrained Matching

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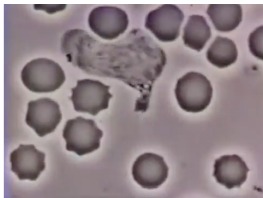
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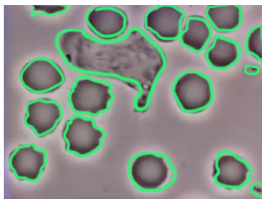


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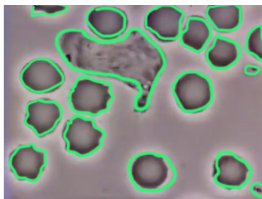


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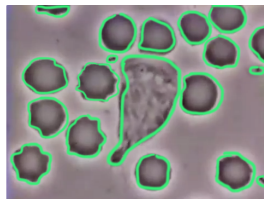
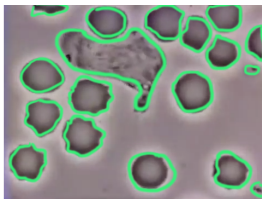


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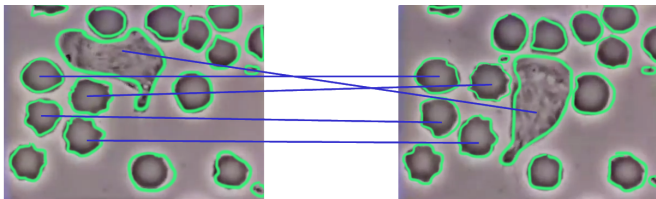


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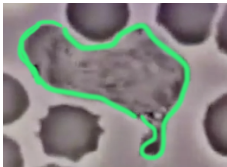
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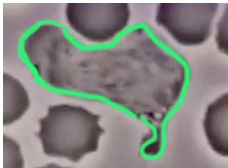


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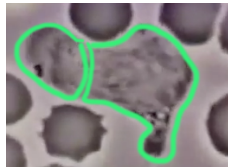
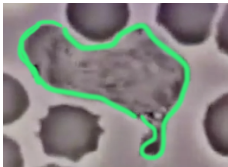


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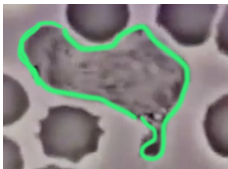


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Challenge: Biological cell division vs. over-segmentation

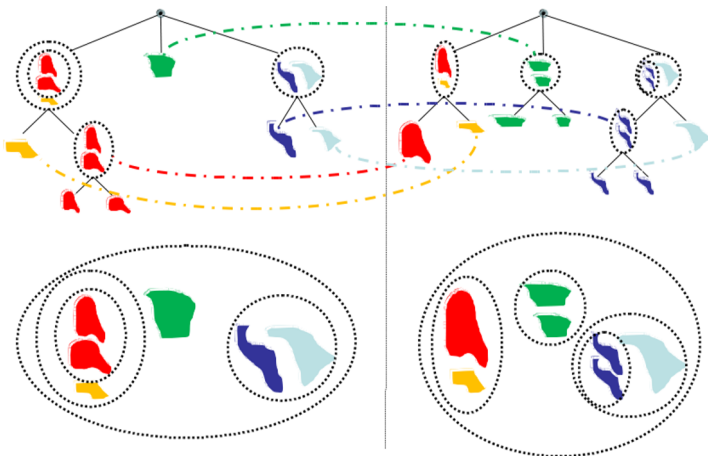
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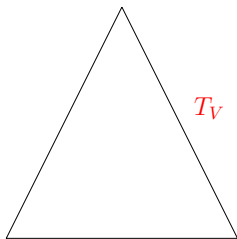
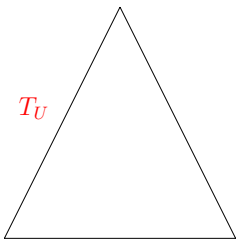
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⇒ [Mosig *et al.*, 2009]: Integrate identification and tracking steps!

# Cosegmentation

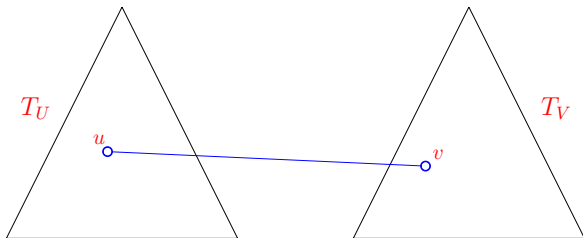


# Tree-Constrained Matching

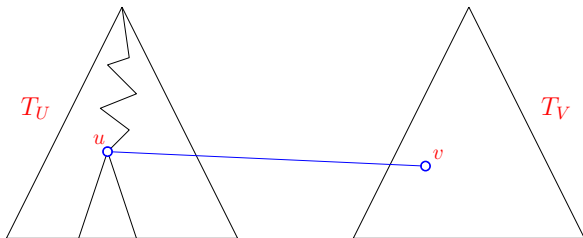




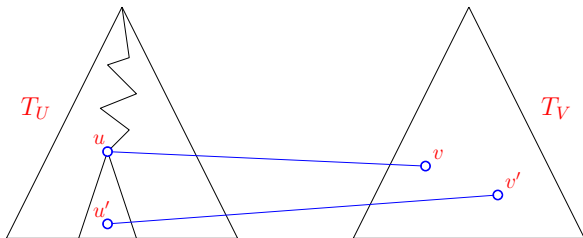
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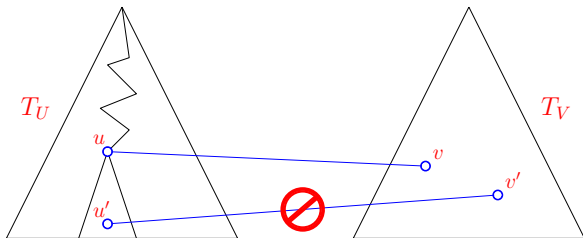
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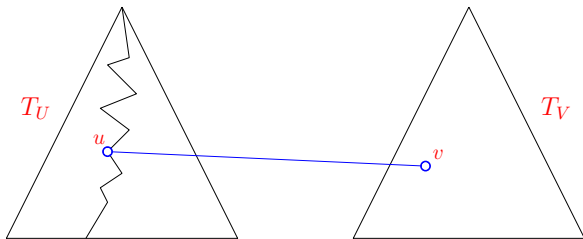
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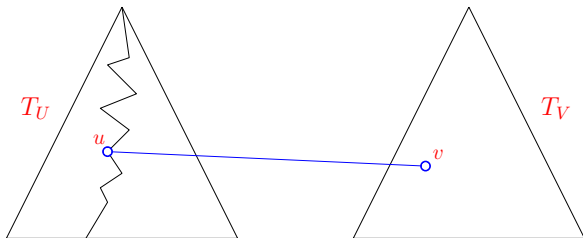
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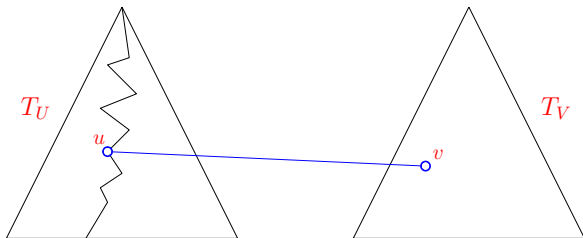


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Given a weighted bipartite graph  $(U, V, E)$  and trees  $T_U$  and  $T_V$  over  $U$  and  $V$ , we want maximum weight matching  $\mathcal{M}$  such that matched vertices in  $T_U$  and  $T_V$  are not comparable

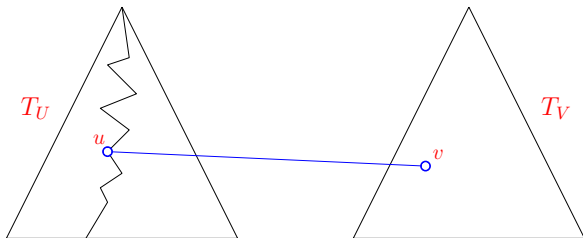
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Unfortunately, as we shall see, this is not the case





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Maximum weight independent set (MIS) in  $d$ -interval graphs:

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Thus, there is a 4-approximation for TCM



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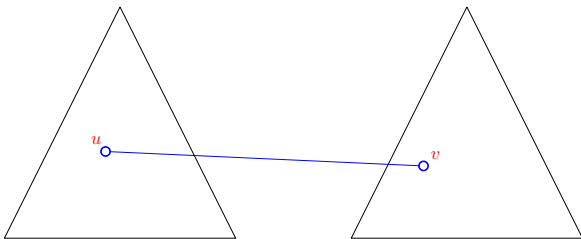
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Generalization to posets

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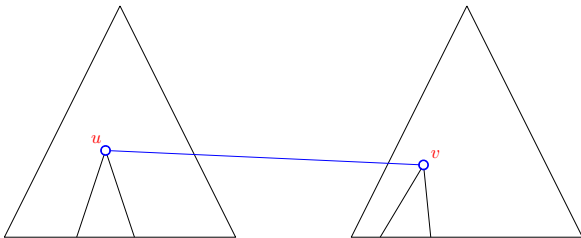
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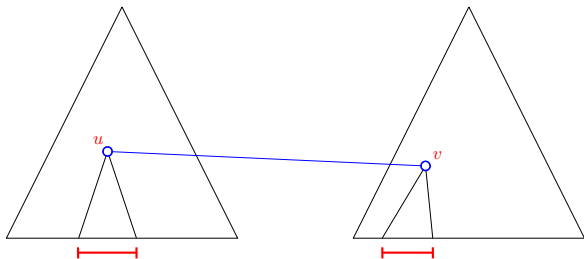
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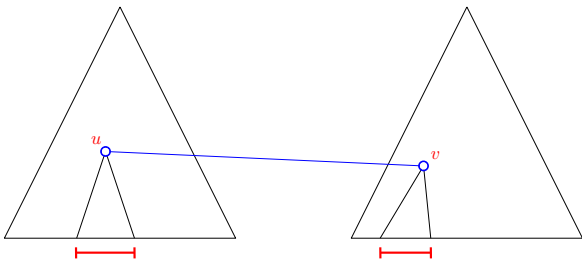
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Matching is feasible  $\Leftrightarrow$  corresp. set of 2-intervals is independent

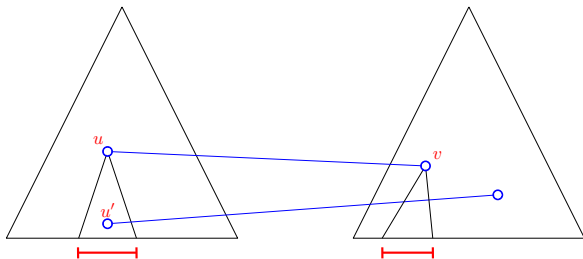




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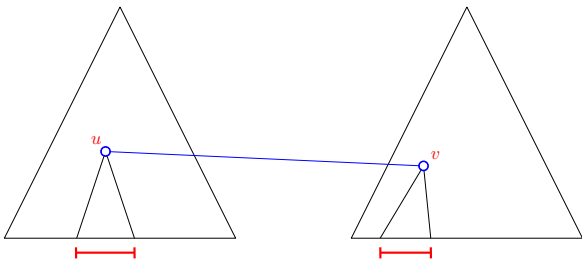
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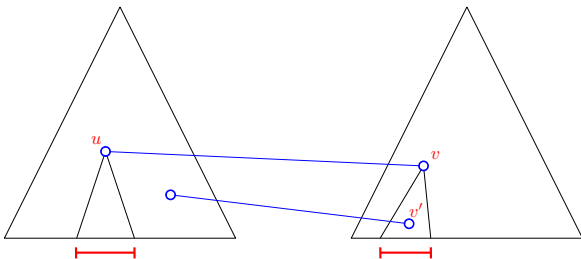




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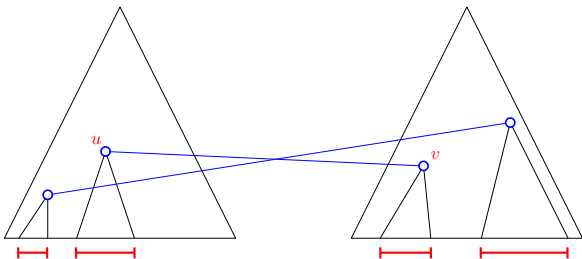




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# Bar-Yehuda *et al.* algorithm

$$\max \sum_{u \in V} x_u$$

$$\text{s.t. } \sum_{u: p \in u} x_u \leq 1 \quad \forall \text{ point } p$$

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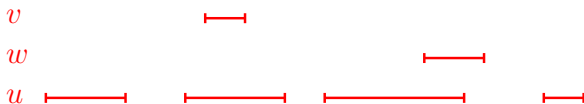
MIS- $d$ -interval( $G$ )

- 1: let  $x$  be optimal LP solution
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- 4:   let  $u$  minimize  $x(N(u))$
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- 6:   remove  $N(u) + u$  from  $G$
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$\forall$  **feasible**  $x$ :  $\exists u : x(N(u)) \leq 2d$

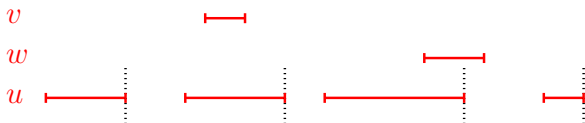
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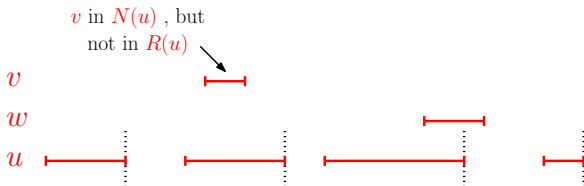
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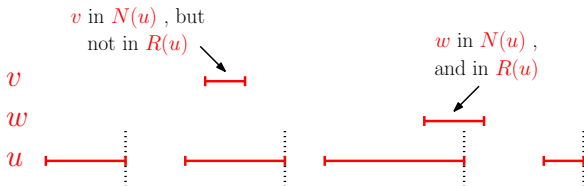
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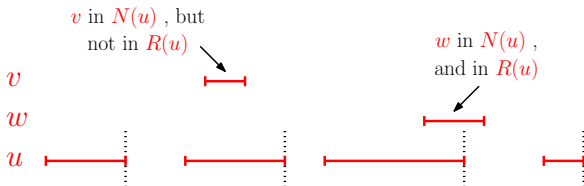
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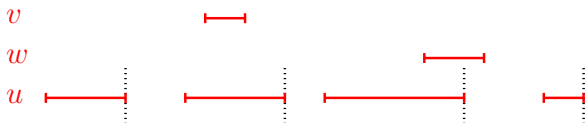






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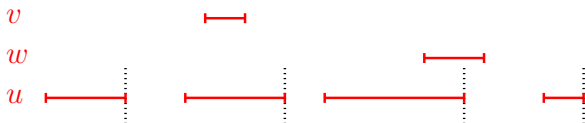
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## 4-approximation for TCM

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$N(e)$  is the set of  
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For ease of analysis, assume that

- For all edges  $x_e > 0$
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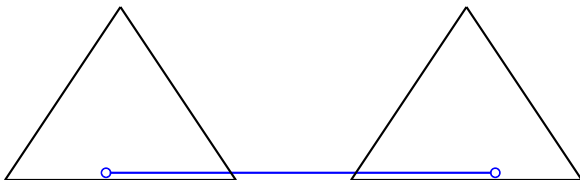
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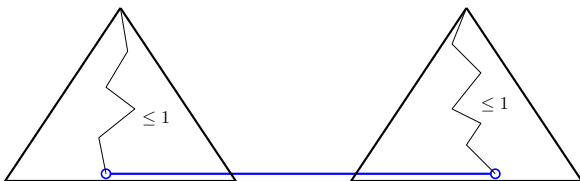


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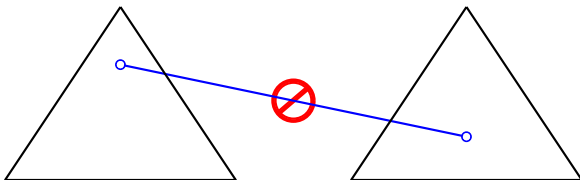
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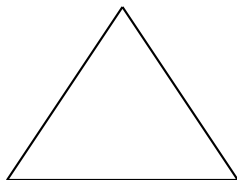
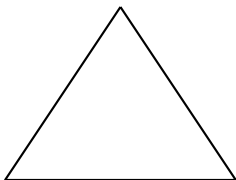
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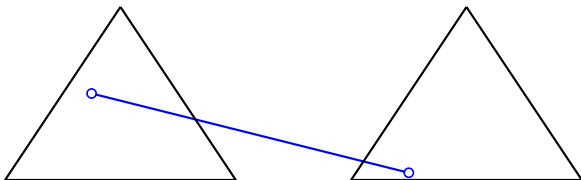
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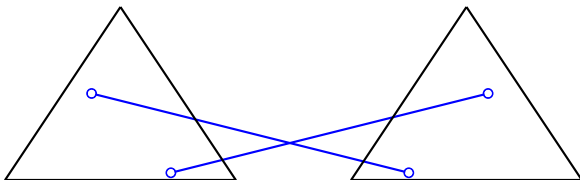
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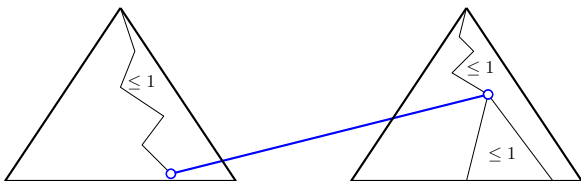
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If we have a leaf-to-leaf edge, we are done

Otherwise, no internal-to-internal edges





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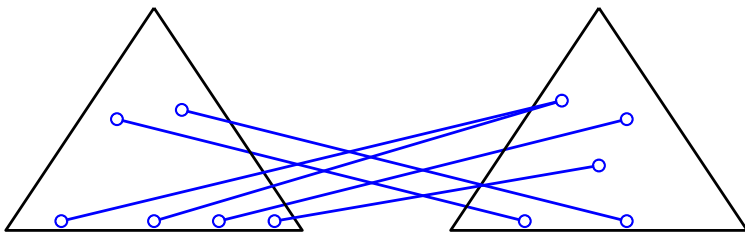
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Integrality gap of LP formulation is  $2 - o(1)$



## 2-approximation

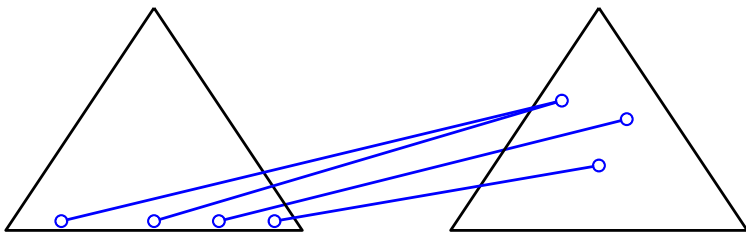
Idea: Exploit bfs structure if  $\forall e : x(N(e)) > 2$





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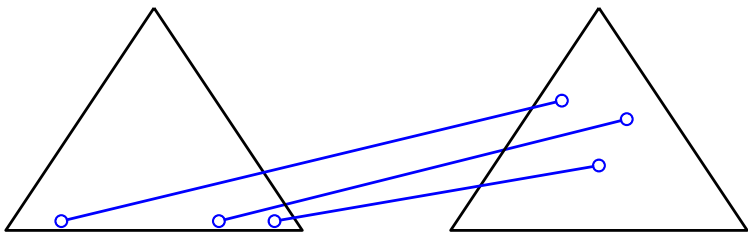
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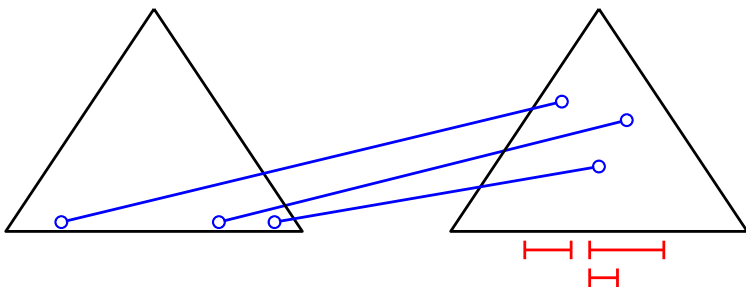
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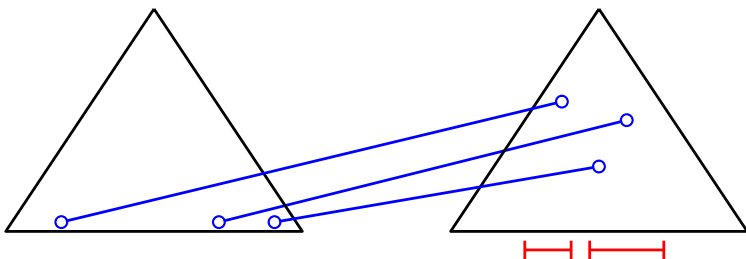
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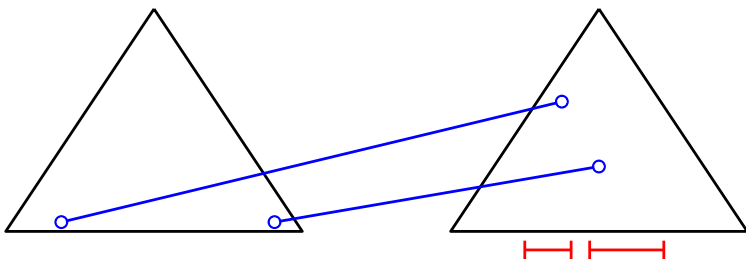






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It cannot be approximated to  $2^{\log^{1-\epsilon} \rho}$ , for any  $\epsilon > 0$ , unless  $NP \subseteq DTIME(n^{\text{poly} \log n})$



**Thank you  
for your  
attention!**