

FASTER TREE EDIT DISTANCE VIA APSP EQUIVALENCE

DIMACS WORKSHOP ON FINE-GRAINED COMPLEXITY OF STRING PROBLEMS

Jakob Nogler⁴

Based on joint work with:

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Virginia Vassilevska Williams⁴ **Yinzhan Xu**³ **Christopher Ye**³

¹ETH Zurich

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³UC San Diego

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(String) Edit Distance Problem

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“Unweighted” (String) Edit Distance: all costs are one

ALGORITHMS FOR (STRING) EDIT DISTANCE

References	Time	Remarks
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MP80	$\mathcal{O}(n^2 / \log^2 n)$	small alphabets and integer weights only
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BI18	$n^{2-o(1)}$	Lower bound under OVH/SETH, already for unweighted

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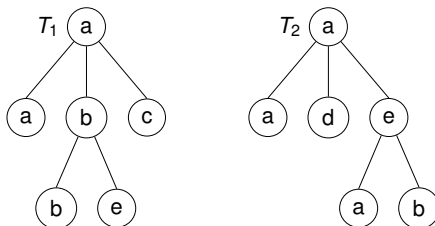
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⇓ Generalization on Trees

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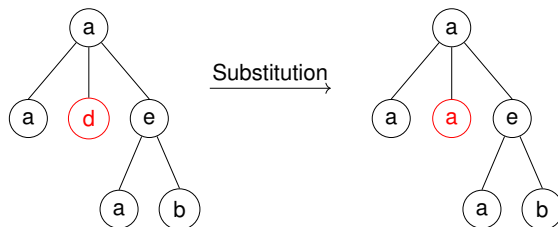
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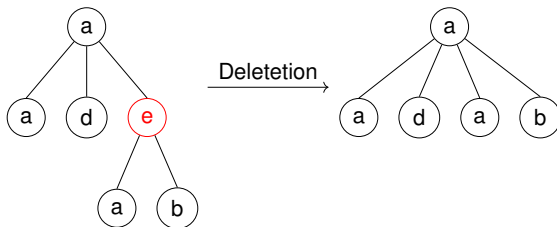
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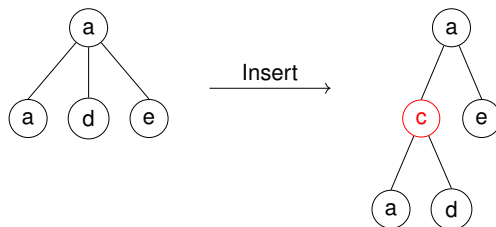
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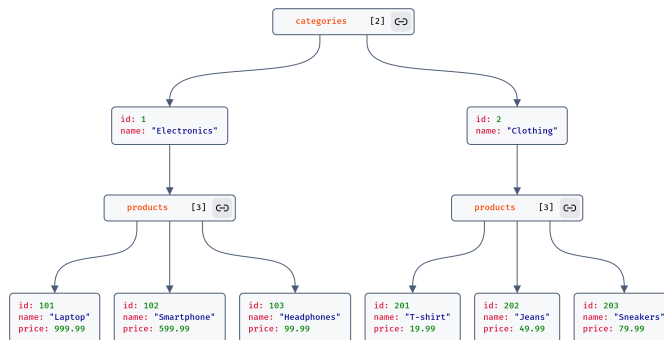


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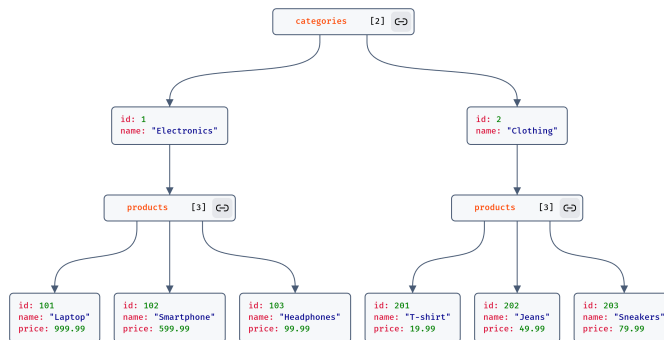
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- ▶ Applications in computational biology, structured data analysis, image processing, compiler optimization, and more.

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Input: A weighted and directed graph G .

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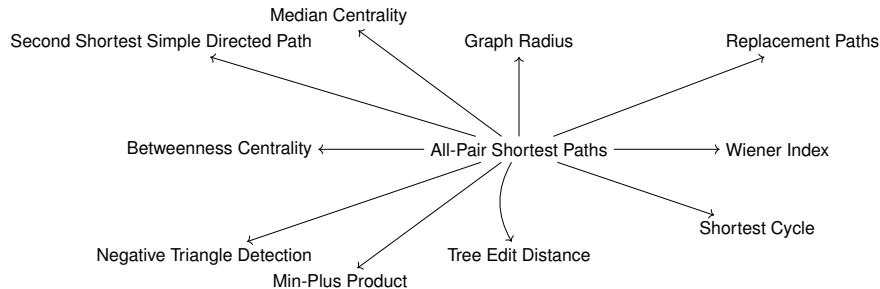
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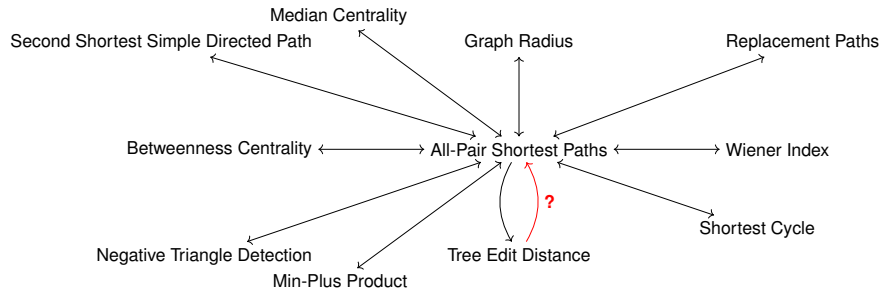
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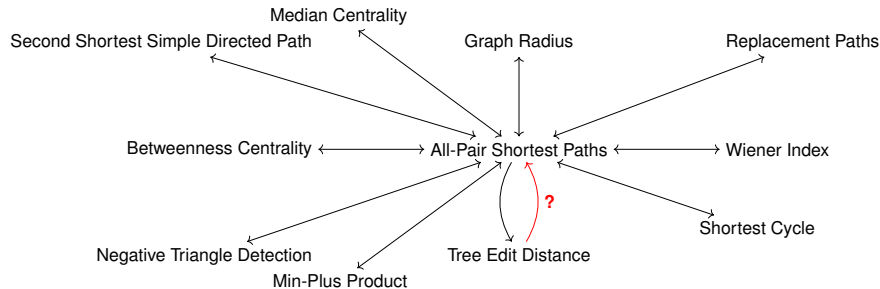
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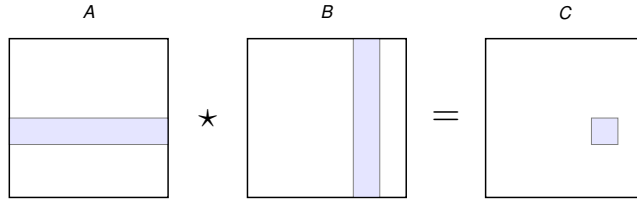


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Question 2: is weighted TED (subcubically) equivalent to APSP?

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Min-plus Product

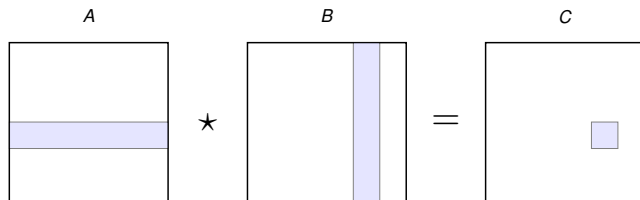


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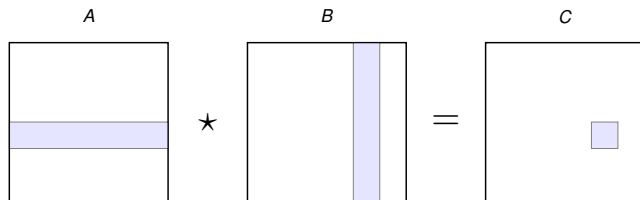
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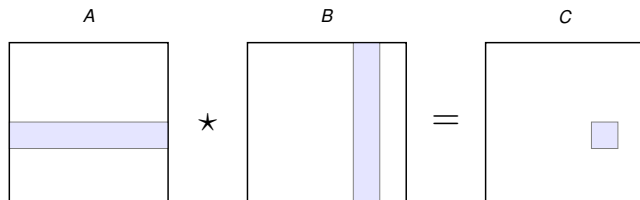
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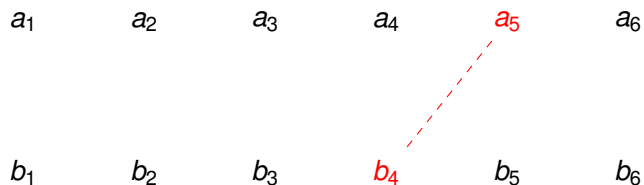
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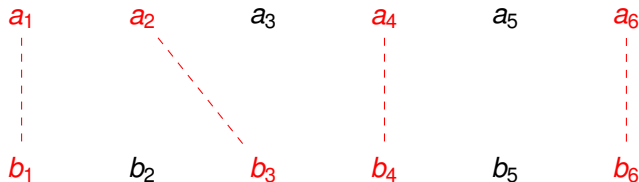


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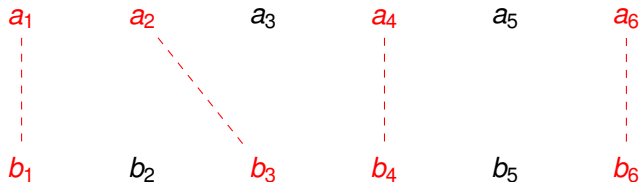
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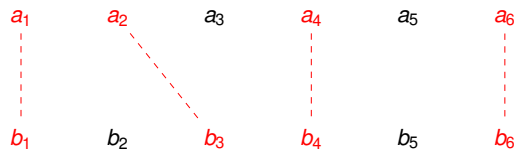
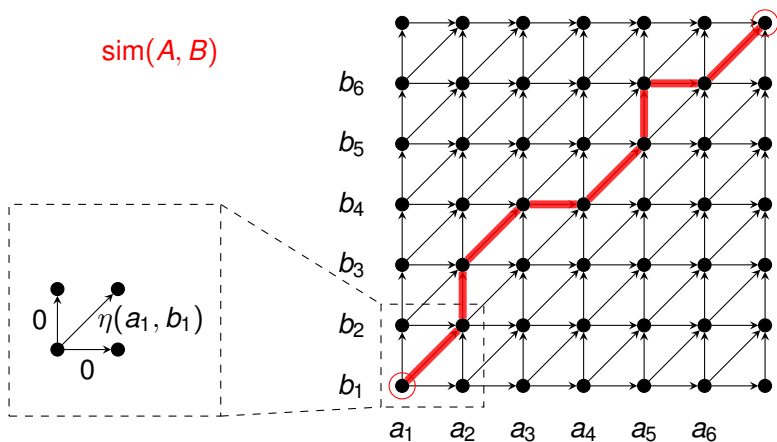
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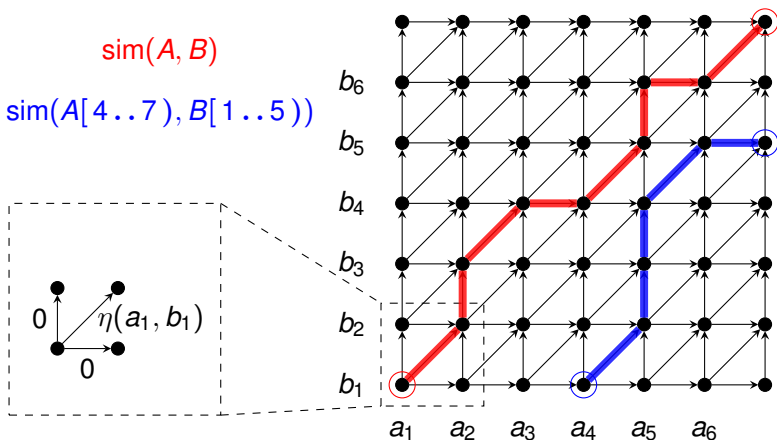
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The **string alignment graph** summarizes the DP scheme computing the similarity.



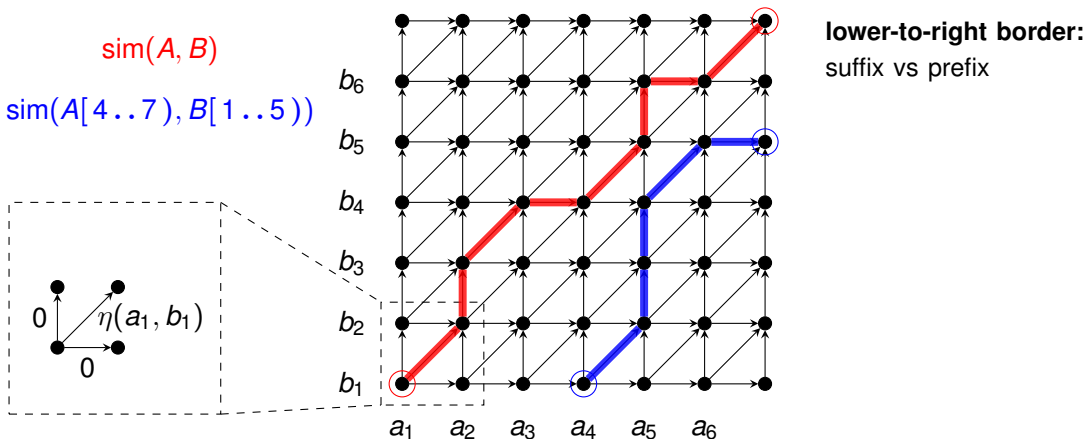
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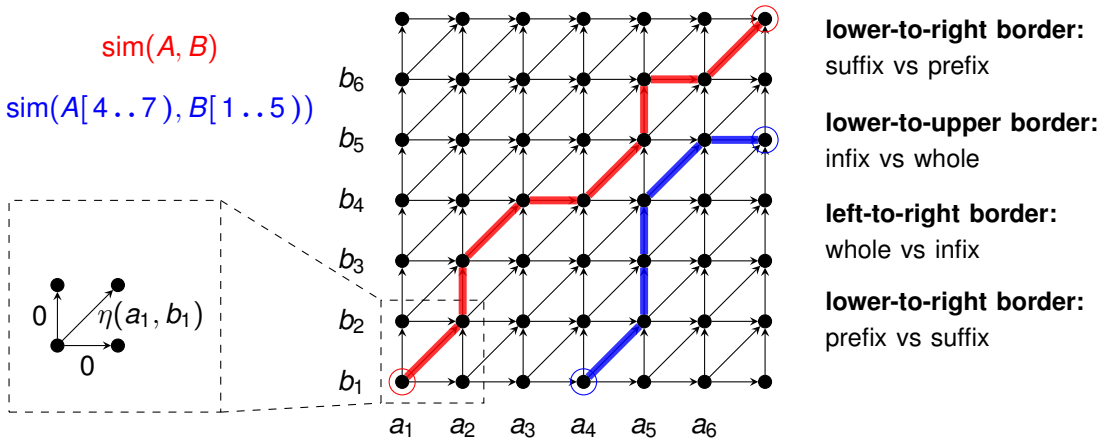
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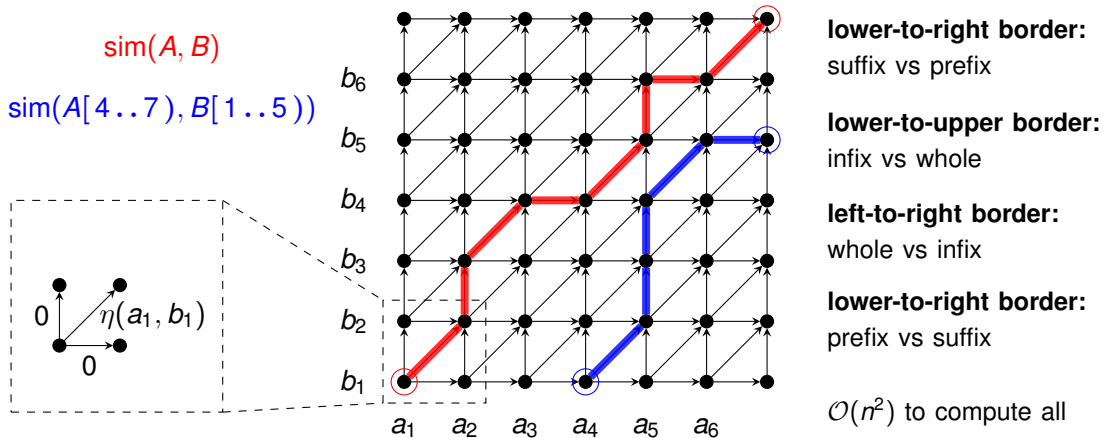
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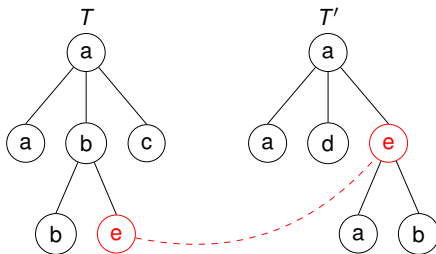
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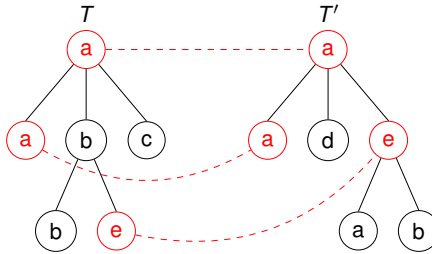


1. $\eta(v, v') := \delta(v, \varepsilon) + \delta(\varepsilon, v') - \delta(v, v')$

How much I save by substituting v with v'

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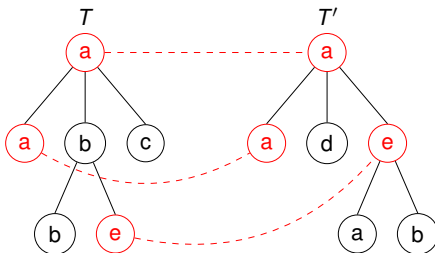
How much I save by substituting v with v'

2. $\text{sim}(T, T') = \text{"maximum weight of similarity matching"}$

Maximum I can save

Goal: Compute the **similarity** of two trees T and T' via a **Min-Plus Product** algorithm.

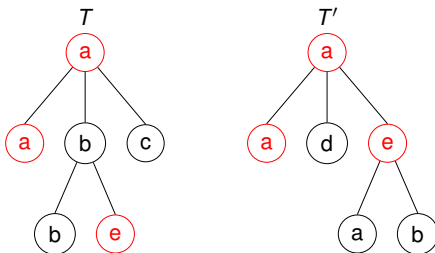
Given two trees T and T' , we define their **similarity** as:



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2. $\text{sim}(T, T') = \text{"maximum weight of similarity matching"}$ *Maximum I can save*
3. Condition on similarity matching: for any two matched vertices (v, v') and (u, u')
 - v is an ancestor of u in T if and only if v' is an ancestor of u' in T' ,
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Goal: Compute the **similarity** of two trees T and T' via a **Min-Plus Product** algorithm.

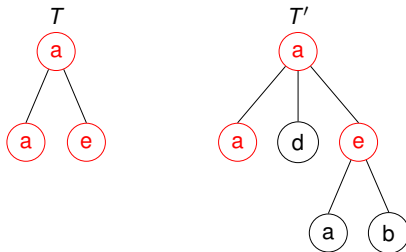
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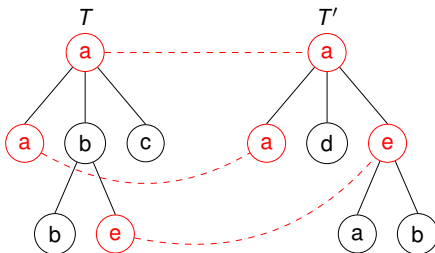
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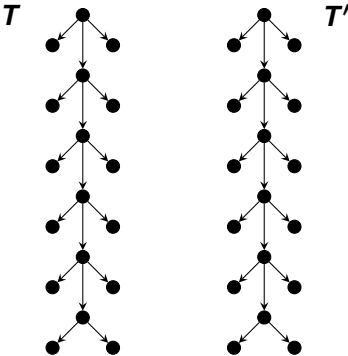
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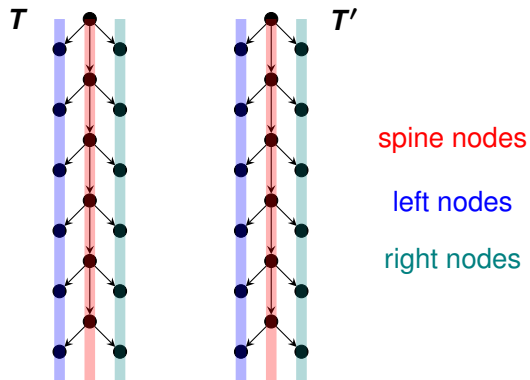


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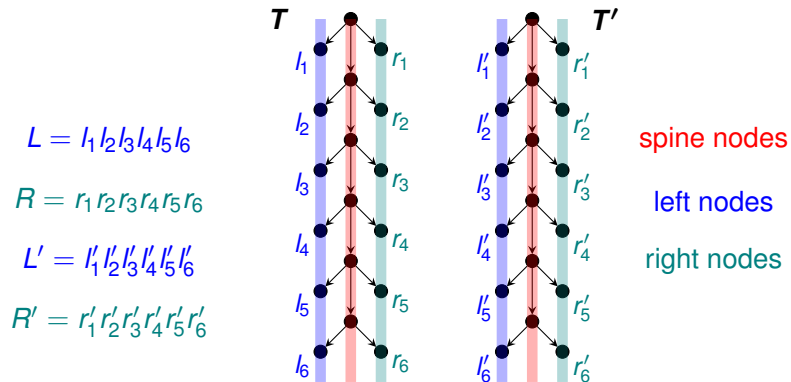
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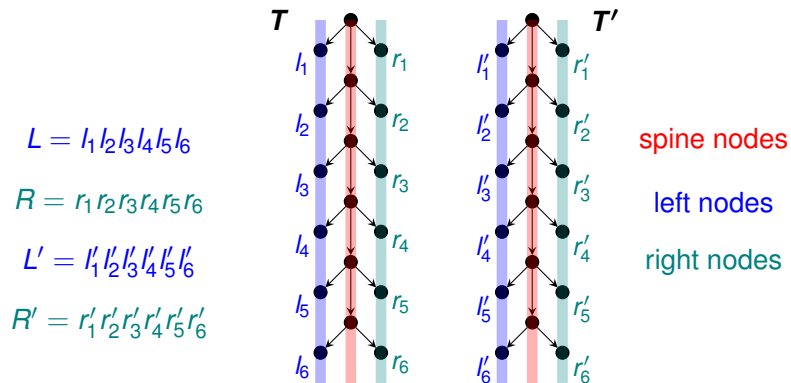
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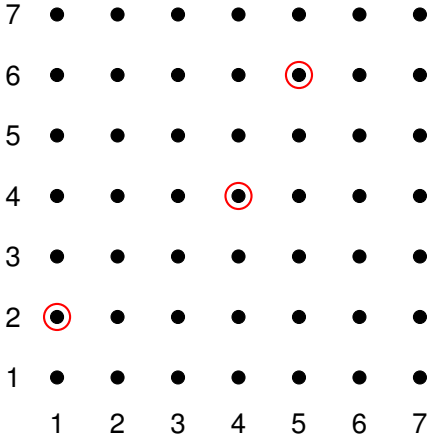
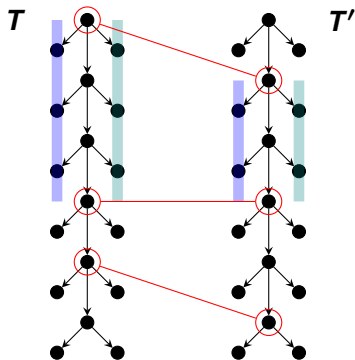
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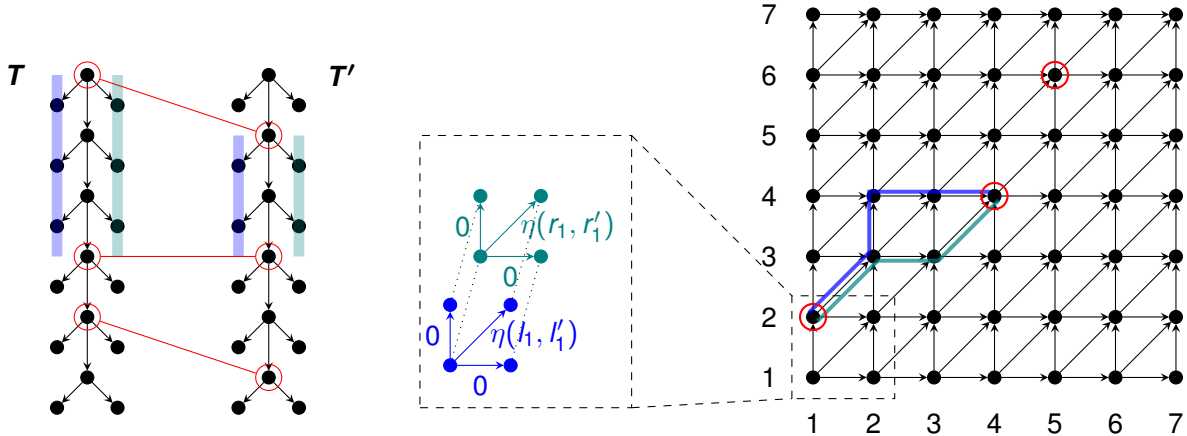
Left-to-Left/Spine-to-spine/Right-to-Right restriction:

spine, **left** and **right** nodes of T only match with nodes of their same type (color) in T' .

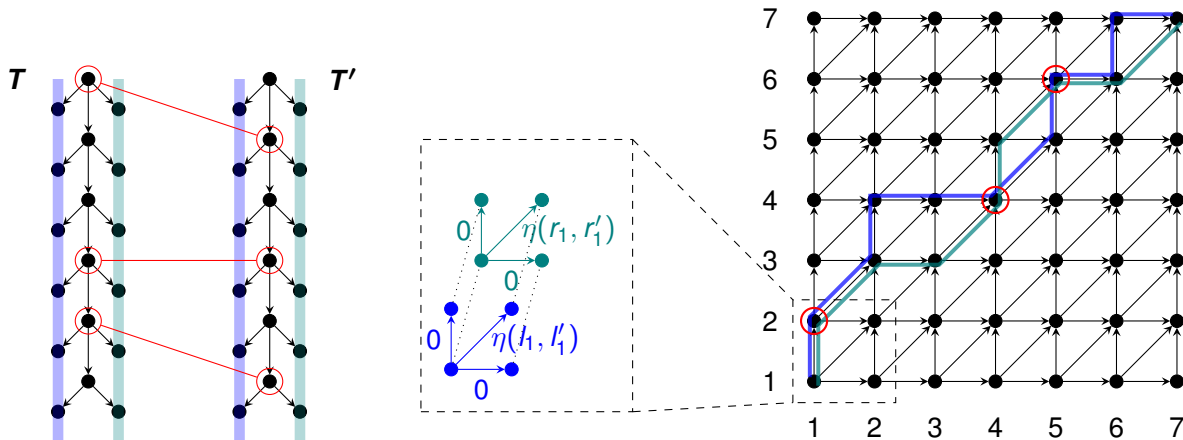
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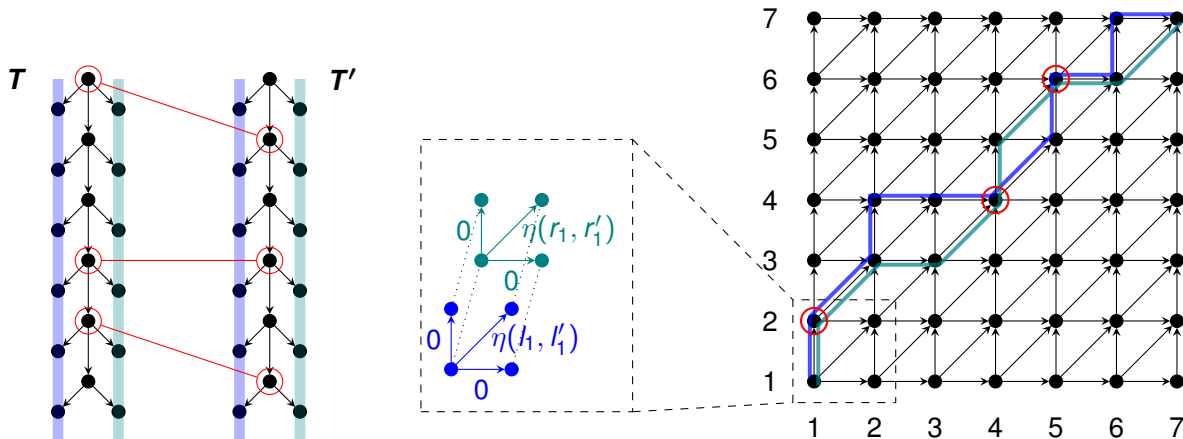
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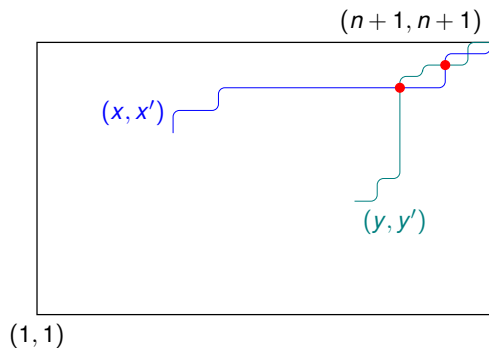
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$\text{sim}(T, T')$ equals to the maximum achievable sum of:

1. the weight of a **path** from $(1, 1)$ to $(n + 1, n + 1)$ in the alignment graph of $\text{sim}(L, L')$;
2. the weight of a **path** from $(1, 1)$ to $(n + 1, n + 1)$ in the alignment graph of $\text{sim}(R, R')$; and
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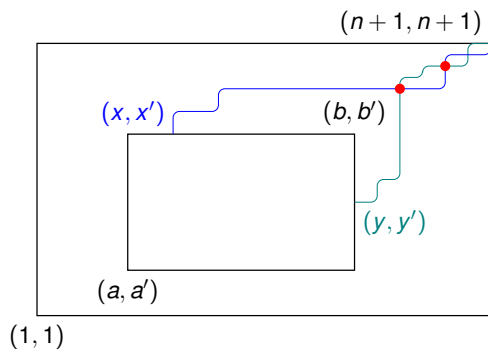
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$\text{sim}((x, x'), (y, y'))$ equals to the maximum achievable sum of:

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Divide et Conquer Scheme

Input:

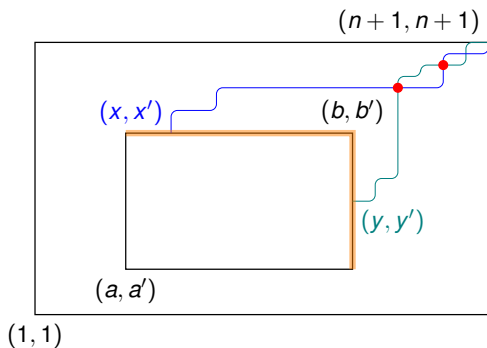
- The lower-left corner (a, a') and upper-right corner (b, b') of a rectangle.

Output:

$\text{sim}((x, x'), (y, y'))$ equals to the maximum achievable sum of:

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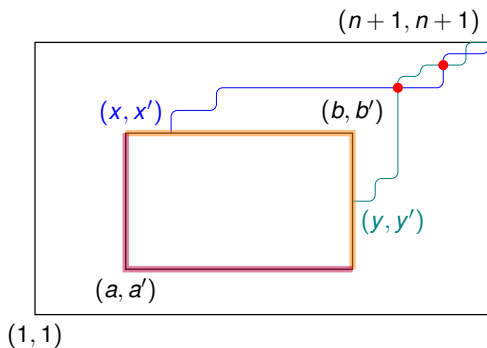
- The lower-left corner (a, a') and upper-right corner (b, b') of a rectangle.
- $\text{sim}((x, x'), (y, y'))$
 $\forall (x, x'), (y, y') \in ([a \dots b] \times \{b'\}) \cup (\{b\} \times [a' \dots b'])$.

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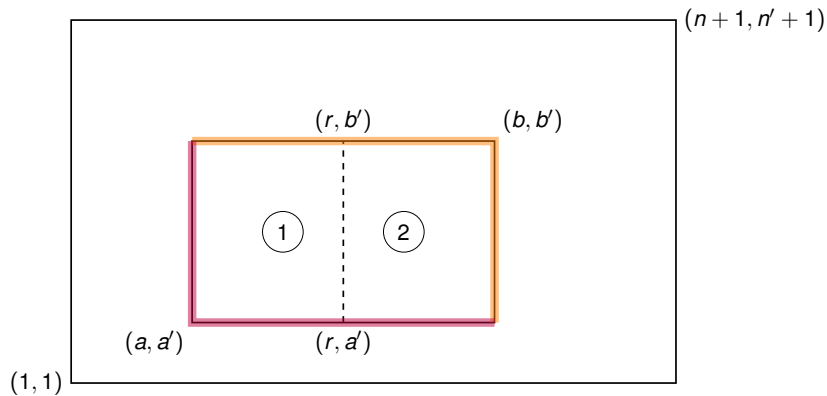
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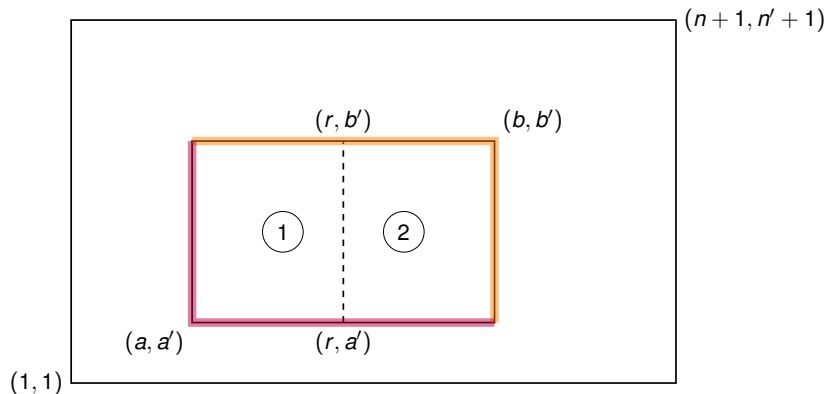
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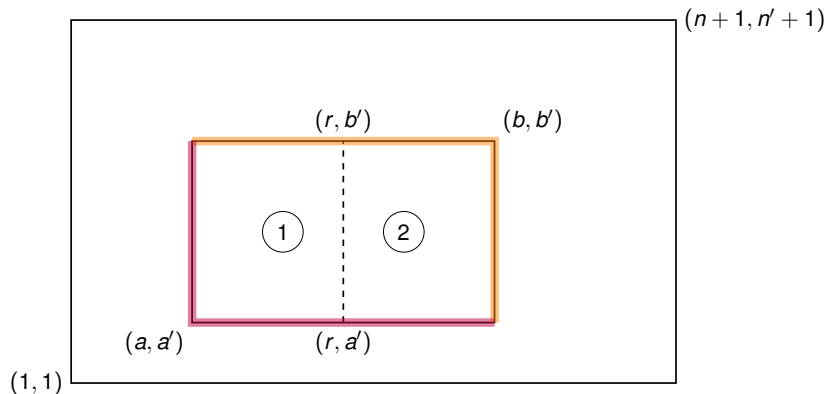
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Inputs of subrectangle 2. ✓

Recurse on subrectangle 2. ✓

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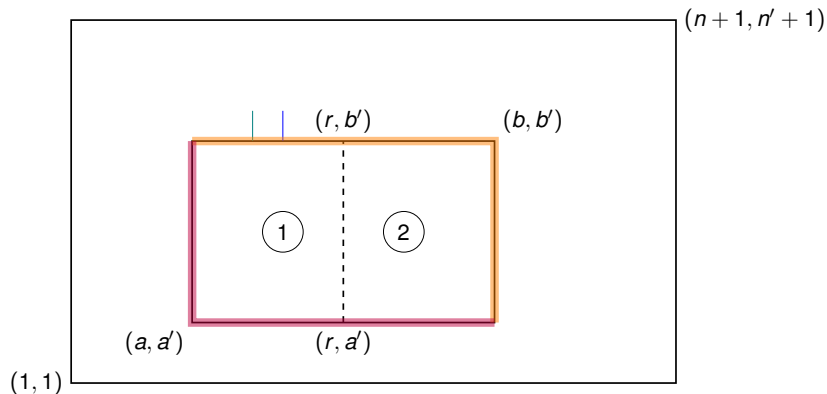


Inputs of subrectangle 2. ✓

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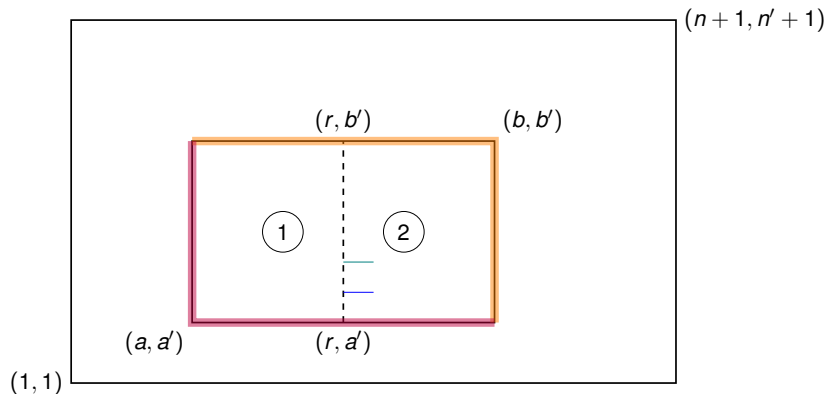


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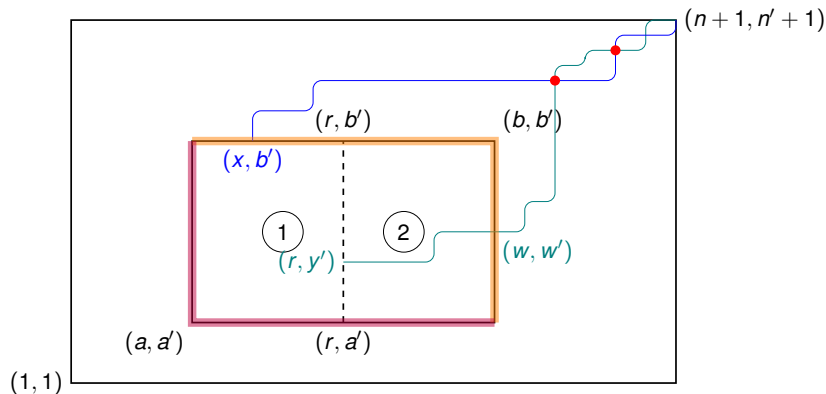


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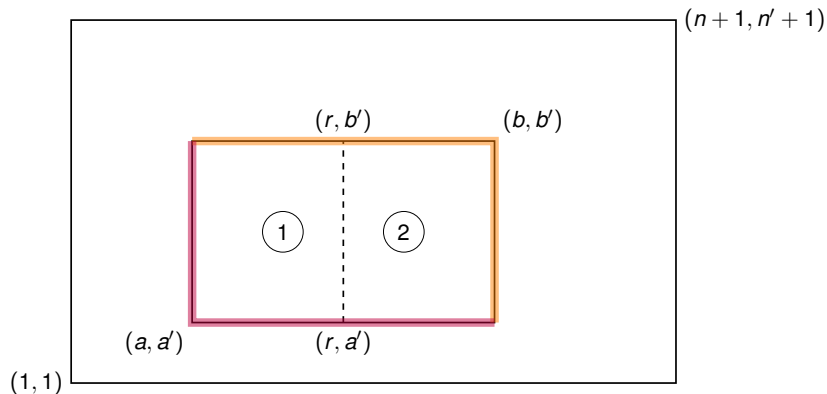
Recurse on subrectangle 2. ✓

Inputs of subrectangle 1. ✓

Recurse on subrectangle 1. ✓

$$\text{sim}((x, b'), (r, y')) = \max_{(w, w')} \left\{ \text{sim}(R[r..w], R'[y'..w']) + \text{sim}((x, b'), (w, w')) \right\}.$$

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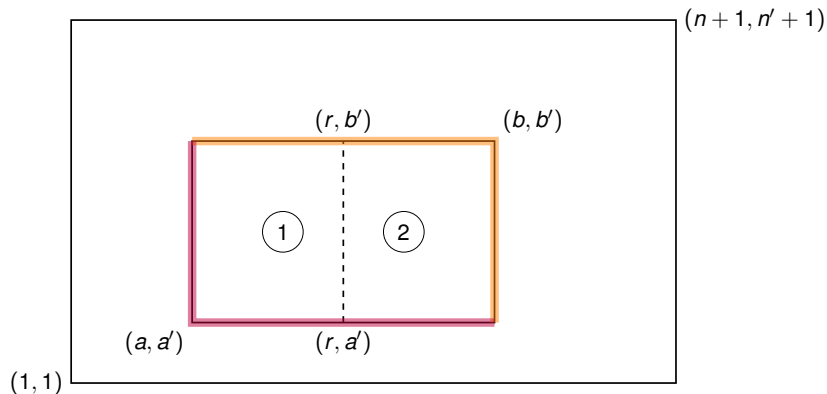
Recurse on subrectangle 2. ✓

Inputs of subrectangle 1. ✓

Recurse on subrectangle 1. ✓

Outputs of big rectangle.

Goal: Compute the **similarity** of two **caterpillar** trees T and T' via a **Min-Plus Product** algorithm **under the Left-to-Left/Spine-to-spine/Right-to-Right restriction**.



Inputs of subrectangle 2. ✓

Recurse on subrectangle 2. ✓

Inputs of subrectangle 1. ✓

Recurse on subrectangle 1. ✓

Outputs of big rectangle. ✓

Goal: Compute the similarity of two caterpillar trees T and T' via a Min-Plus Product algorithm under the Left-to-Left/Spine-to-spine/Right-to-Right restriction.

- ▶ Since we can handle vertical “cuts”, we can also handle horizontal “cuts”.



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- ▶ We obtain the recurrence

$$T(n) = 4T(n/2) + \mathcal{O}(T_{\text{APSP}}).$$

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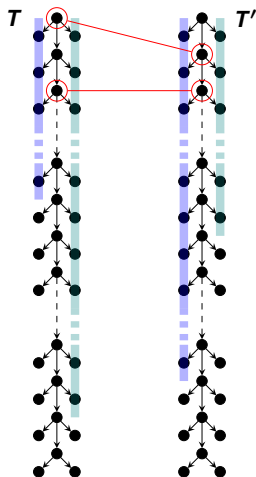
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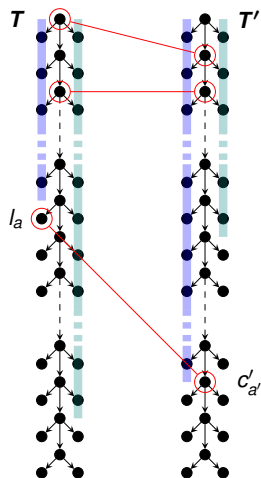
Thus, $T(n) = \mathcal{O}(T_{\text{APSP}})$ assuming $T_{\text{APSP}} = \mathcal{O}(n^{2+\varepsilon})$ for some $\varepsilon > 0$.

- ▶ I cheated a bit... in the scheme I need to remember whether spines nodes are already mapped.

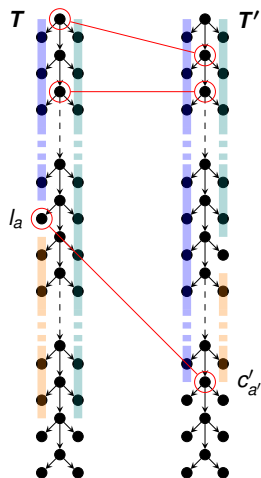
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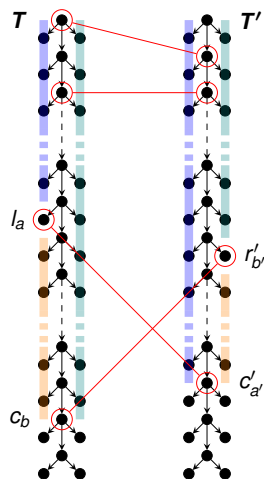
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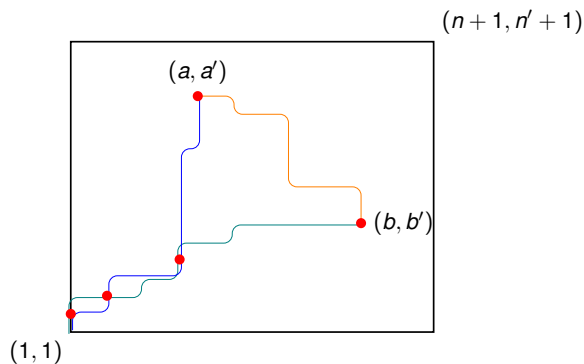
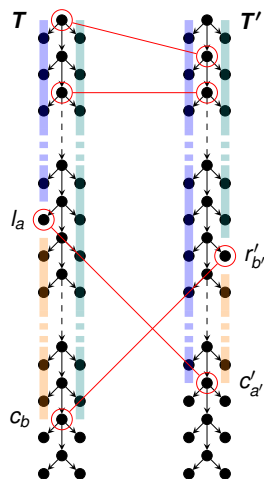
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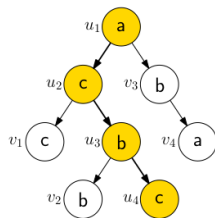
We generalize similarity computation of caterpillar trees to Spine Edit Distance.

Goal: Compute the **similarity** of two **caterpillar** trees T and T' via a **Min-Plus Product** algorithm.

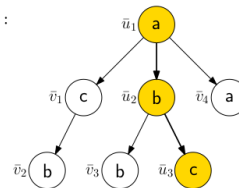
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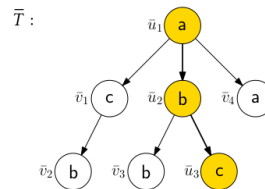
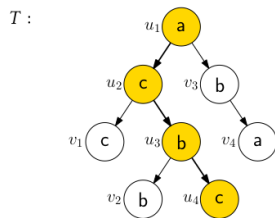
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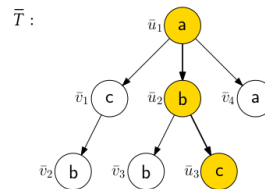
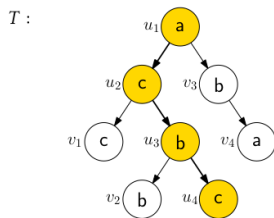


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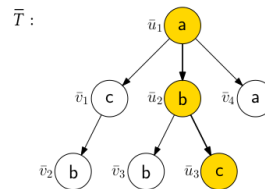
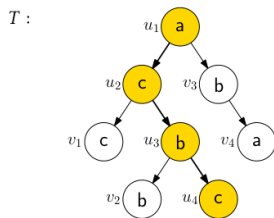
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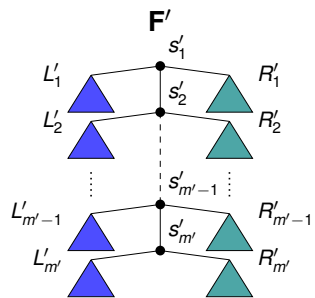
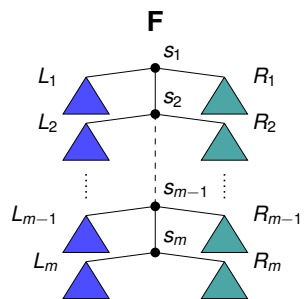


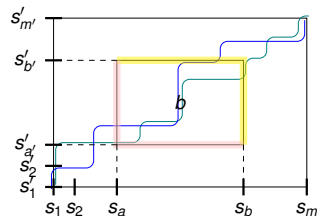
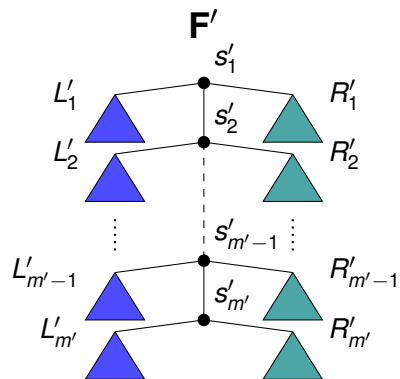
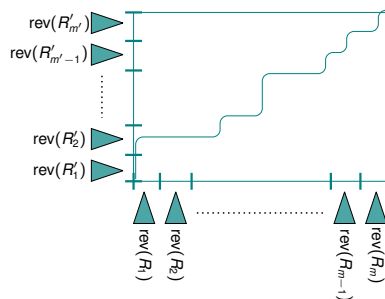
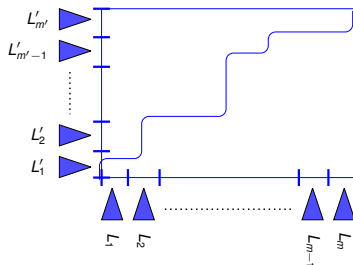
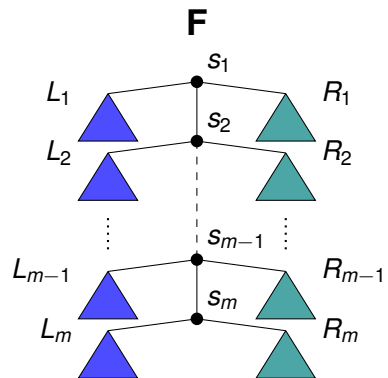
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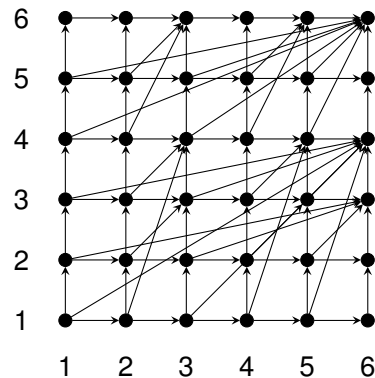
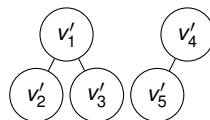
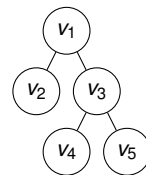
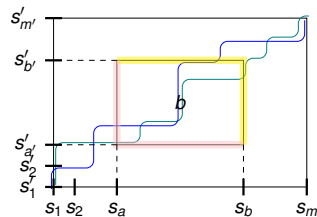
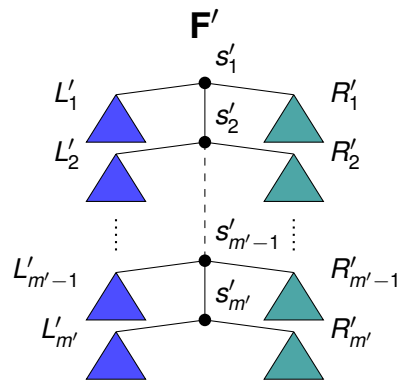
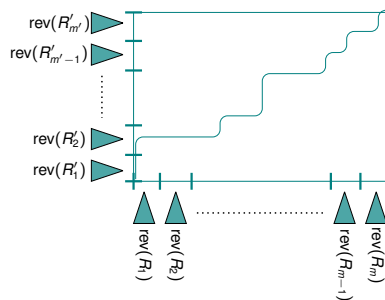
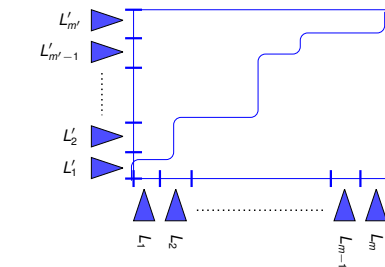
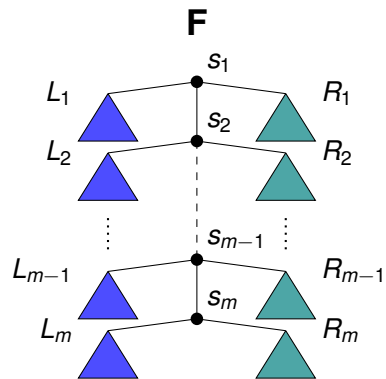
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We also prove that similarity computation on arbitrary trees is fine-grained equivalent to Spine Edit Distance.







TWO OPEN PROBLEMS

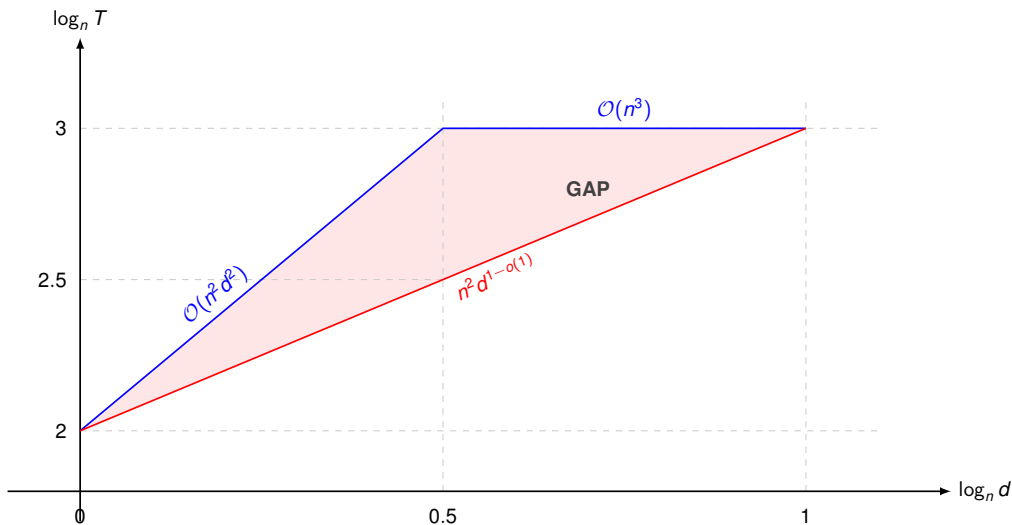
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[Charalampopoulos, Gawrychowski, Mozes and Weimann, SPIRE 2022]

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Thanks!