# Similarity Search The Tree Edit Distance

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## Outline

- Tree Edit Distance
  - Preliminaries and Definition
  - Forests Distance and Recursive Formula
  - Second Recursive Formula
  - The Tree Edit Distance Algorithm
  - Example: Tree Edit Distance Computation
  - Complexity of the Tree Edit Distance Algorithm

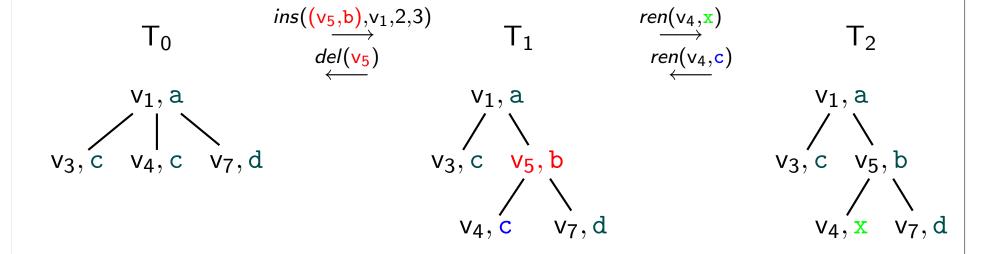
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## **Edit Operations**

- We assume ordered, labeled trees
- Rename node: ren(v, l')
  - change label I of v to  $I' \neq I$
- Delete node: del(v) (v is not the root node)
  - remove v
  - connect v's children directly to v's parent node (preserving order)
- Insert node: ins(v, p, k, m)
  - insert new node v as a child of p at position k
  - substitute children  $c_k, c_{k+1}, \ldots, c_m$  of p with v
  - insert  $c_k, c_{k+1}, \ldots, c_m$  as children of the new node v (preserving order)
- Insert and delete are inverse edit operations (i.e., insert undoes delete and vice versa)

## Example: Edit Operations



#### **Edit Cost Function**

- Represent edit operation as node pair  $(a, b) \neq (\varepsilon, \varepsilon)$  (written also as  $a \rightarrow b$ ,  $\varepsilon$  is the null node)
  - a  $\rightarrow \varepsilon$ : delete a
  - $\varepsilon \rightarrow b$ : insert b
  - a  $\rightarrow$  b: rename a to b
- Cost function  $\alpha(a \rightarrow b)$ :
  - assign to each edit operation a non-negative real
  - cost can be different for different nodes
  - we use constant costs  $\omega_{\textit{ins}}, \omega_{\textit{del}}, \omega_{\textit{ren}}$
- We constrain  $\alpha$  to be a distance metric:
  - (i) triangle inequality:  $\alpha(a \rightarrow b) + \alpha(b \rightarrow c) \ge \alpha(a, c)$
  - (ii) symmetry:  $\alpha(a \rightarrow b) = \alpha(b \rightarrow a)$
  - (iii) identity:  $\alpha(a \rightarrow b) = 0 \Leftrightarrow \lambda(a) = \lambda(b)$

#### Definition

#### Definition (Tree Edit Distance)

The tree edit distance between two trees is the minimum cost sequence of node edit operations (node deletion, node insertion, node rename) that transforms on tree into the other.

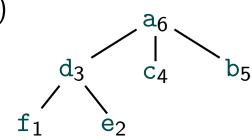
• Cost of a sequence  $S = \{s_1, \dots, s_n\}$  of edit operations:

$$\alpha(S) = \sum_{i=1}^{i=n} \alpha(s_i)$$

• As the cost function is a metric, also the tree edit distance is a metric.

#### Postorder Traversal

- Postorder traversal of an ordered tree:
  - traverse subtrees rooted in children of current node (from left to right) in postorder
  - visit current node
- Example: postorder = (f, e, d, c, b, a)



- Observations: The postorder number of a node is larger than
  - the postorder numbers of all its descendants
  - the postorder numbers of all its left siblings

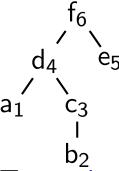
#### Subtrees and Subforests

- A subtree T' of T is a tree that consists of:
  - a subset of the nodes of T:  $N(T') \subseteq N(T)$
  - all edges in T that connect these nodes:  $E(T') \subseteq E(T)$
- Ordered Forests:
  - a forest is a set of trees
  - an ordered forest is a sequence of trees
- Ordered Subforests of a tree T:
  - formed by subtrees of T with disjoined nodes
  - subtrees ordered by the postorder number in T of their root

## Example: Subtrees and Subforests

• Example tree (postorder numbers are node IDs):

$$T = (\{v_1, v_2, v_3, v_4, v_5, v_6\}, \{(v_6, v_4), (v_6, v_5), (v_4, v_1), (v_4, v_3), (v_3, v_2)\})$$



• Two subtrees of T:

$$T_1' = (\{v_3\}, \{\}) \qquad T_2' = (\{v_4, v_1, v_3\}, \{(v_4, v_1), (v_4, v_3)\})$$

$$c_3 \qquad d_4 \qquad c_3$$

Ordered subforest of T:

$$F = ((\{v_2\}, \{\}), (\{v_4, v_1, v_3\}, \{(v_4, v_1), (v_4, v_3)\}), (\{v_5\}, \{\}))$$

$$b_2$$
  $d_4$   $e_5$   $a_1$   $c_3$ 

## Notation I/II

- We use the following notation:
  - T[i] is the i-th node of T in postorder (we say: T[i] is node i of T)
  - T[i..j] is the subforest formed by the nodes T[i] to T[j]
  - I(i) is the left-most leaf descendant of node T[i]
  - desc(T[i]) is the set of all descendants of T[i] including T[i] itself (elements of desc(T[i]) are usually denoted with  $d_i$ )
- Node identifiers:
  - we assume that the node IDs correspond to their postorder number
  - we refer to a node simply by its ID, if the context is clear

## Notation II/II

- T[I(i)..i] is the subtree rooted in T[i], i.e., the subtree consisting of node i and all its descendants
- A special subforests of the form

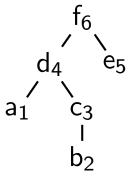
$$T[I(i)..d_i], \qquad (d_i \in desc(T[i]))$$

is a prefix of the subtree rooted in T[i].

- Observations:
  - If a node k is in  $T[I(i)..d_i]$ , also all its descendants are in  $T[I(i)..d_i]$ .
  - A (sub)tree with n nodes has n prefixes.

## Example: Subtrees and Subforests

• Example tree:



- Descendants:  $desc(T[4]) = \{T[1], T[2], T[3], T[4]\}$
- Left-most leaf descendants: I(1) = I(4) = I(6) = T[1]
- Some ordered subforests of the form  $T[I(i)..d_i]$ ,  $d_i \in desc(i)$ :

T[/(4)3]	T[/(4)4]	T[/(6)5]	T[/(5)5]
a <sub>1</sub> c <sub>3</sub> I b <sub>2</sub>	$d_4$ $a_1$ $c_3$ $b_2$	$d_4$ $e_5$ $a_1$ $c_3$ $b_2$	e <sub>5</sub>

## Edit Mapping

#### Definition (Edit Mapping)

An edit mapping M between  $T_1$  and  $T_2$  is a set of node pairs that satisfy the following conditions:

- (1)  $(a,b) \in M \Rightarrow a \in N(T_1), b \in N(T_2)$
- (2) for any two pairs (a, b) and (x, y) of M:
  - (i)  $a = x \Leftrightarrow b = y$  (one-to-one condition)
  - (ii) a is to the left of  $x^1 \Leftrightarrow b$  is to the left of y (order condition)
  - (iii) a is an ancestor of x ⇔ b is an ancestor of y (ancestor condition)
- (3) Optional:  $a = root(T_1)$  and  $b = root(T_1) \Rightarrow (a, b) \in M$  (forbid deleting the root node)

<sup>&</sup>lt;sup>1</sup>i.e., a precedes x in both preorder and postorder

## Edit Mapping

The cost of the mapping is

$$\alpha(M) = \sum_{(a,b)\in M} \alpha(a \to b) + \sum_{a\in D} \alpha(a \to \varepsilon) + \sum_{b\in I} \alpha(\varepsilon \to b),$$

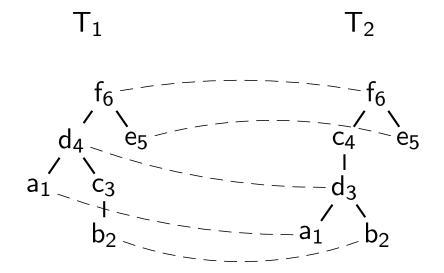
where D and I are the nodes of  $T_1$  and  $T_2$ , respectively, not touched by a line in M.

• Alternative definition of the tree edit distance  $ted(T_1, T_2)$ :

$$ted(T_1, T_2) = min\{\alpha(M) \mid M \text{ is a mapping from } T_1 \text{ to } T_2\}$$

## Example: Mapping

- $M = \{(T_1[6], T_2[6]), (T_1[5], T_2[5]), (T_1[4], T_2[3]), (T_1[1], T_2[1]), (T_1[2], T_2[2])\}$ 
  - T<sub>1</sub>[3] is deleted from T<sub>1</sub>
  - $T_2[4]$  is inserted into  $T_2$
  - no proper rename (only rename to the same label with cost 0)



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#### Forest Distance

#### Definition (Forest Distance)

The forest distance between two ordered forests is the minimum cost sequence of node edit operations (node deletion, node insertion, node rename) that transforms on forest into the other.

- Edit mapping and edit operations in a forest:
  - Each tree in the forest has a root node.
  - We imagine a dummy node that is the parent of all this root nodes.
  - The sibling order in the imaginary tree is the tree order in the forest.
  - The dummy node connects the forest to become a tree.
  - Then all edit operations and edit mappings valid between two imaginary trees are valid also between the respective forests.
- The tree edit distance is a special case of the forest distance, where the forest has the form T[I(i)..i], i.e. it consists of a single tree.

## Recursive Formula: Distance to the Empty Forest

#### Lemma (Empty Forest [ZS89, AG97])

Given two trees  $T_1$  and  $T_2$ ,  $i \in N(T_1)$  and  $d_i \in desc(i)$ ,  $j \in N(T_2)$  and  $d_j \in desc(j)$ , then:

- (i)  $fdist(\emptyset, \emptyset) = 0$
- (ii)  $fdist(\mathsf{T}_1[I(i)..\mathsf{d}_i],\emptyset) = fdist(\mathsf{T}_1[I(i)..\mathsf{d}_i-1],\emptyset) + \omega_{del}$
- (iii)  $fdist(\emptyset, T_2[I(j)..d_j]) = fdist(\emptyset, T_2[I(j)..d_j 1]) + \omega_{ins}$

#### Proof.

Case (i) requires no edit operation. In cases (ii), the distance corresponds to the cost of deleting all nodes in  $T_1[I(i)..d_i]$ . In cases (iii), the distance corresponds to the cost of inserting all nodes in  $T_2[I(j)..d_i]$ .

#### First Recursive Formula: Forest Distance

#### Lemma (First Recursive Formula)

Given two trees  $T_1$  and  $T_2$ ,  $i \in N(T_1)$  and  $d_i \in desc(i)$ ,  $j \in N(T_2)$  and  $d_i \in desc(j)$ , then:

$$fdist(\mathsf{T}_1[I(i)..\mathsf{d}_i],\mathsf{T}_2[I(j)..\mathsf{d}_i]) = \min$$

$$fdist(\mathsf{T}_{1}[I(i)..\mathsf{d}_{i}-1],\mathsf{T}_{2}[I(j)..\mathsf{d}_{j}]) + \omega_{del} \\ fdist(\mathsf{T}_{1}[I(i)..\mathsf{d}_{i}],\mathsf{T}_{2}[I(j)..\mathsf{d}_{j}-1]) + \omega_{ins} \\ fdist(\mathsf{T}_{1}[I(i)..I_{(d_{i})}-1],\mathsf{T}_{2}[I(j)..I_{(d_{j})}-1]) \\ + fdist(\mathsf{T}_{1}[I(d_{i})..\mathsf{d}_{i}-1],\mathsf{T}_{2}[I(d_{j})..\mathsf{d}_{j}-1]) \\ + \omega_{ren} \\$$

#### Proof

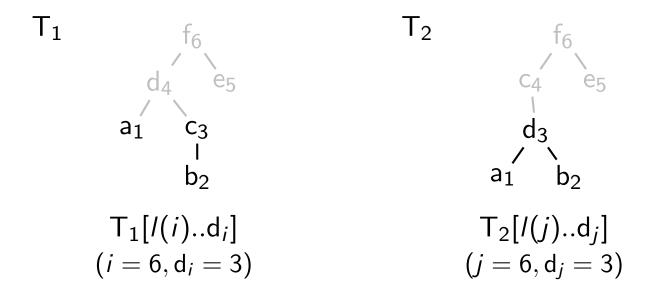
#### Proof.

Let M be the minimum-cost map between  $T_1[I(i)..d_i]$  and  $T_2[I(j)..d_i]$ , i.e., the map we are looking for. Then for  $T_1[d_i]$  and  $T_2[d_i]$  there are tree possibilities:

- (1)  $T_1[d_i]$  is not touched by a line in M:  $T_1[d_i]$  is deleted and  $fdist(T_1[I(i)..d_i], T_2[I(j)..d_i]) = fdist(T_1[I(i)..d_i - 1], T_2[I(j)..d_i]) + \omega_{del}$
- (2)  $T_2[d_i]$  is not touched by a line in M:  $T_2[d_i]$  is inserted and  $fdist(\mathsf{T}_1[I(i)..\mathsf{d}_i],\mathsf{T}_2[I(j)..\mathsf{d}_i]) = fdist(\mathsf{T}_1[I(i)..\mathsf{d}_i],\mathsf{T}_2[I(j)..\mathsf{d}_i-1]) + \omega_{ins}$
- (3) Both,  $T_1[d_i]$  and  $T_2[d_j]$  are touched by a line in M: We show (by contradiction) that in this case  $(T_1[d_i], T_2[d_i]) \in M$ , i.e.,  $T_1[d_i]$  is renamed to  $T_2[d_i]$ : Assume  $(T_1[d_i], T_2[x]) \in M$  and  $(T_1[y], T_2[d_i]) \in M$ .
  - Case  $T_1[d_i]$  is to the right of  $T_1[y]$ : By sibling condition on M also  $T_2[d_i]$  must be to the right of  $T_2[x]$ . Impossible in  $T_2[I(j)..d_i]$ .
  - Case  $T_1[d_i]$  is proper ancestor of  $T_1[y]$ : By ancestor condition on M also  $T_2[d_i]$  must be ancestor of  $T_2[x]$ . Impossible in  $T_2[I(j)..d_i]$ .

As these three cases express all possible mappings yielding  $fdist(T_1[I(i)..d_i], T_2[I(j)..d_i])$ , we take the minimum of these tree costs.

## Example: First Recursive Formula (1/3)

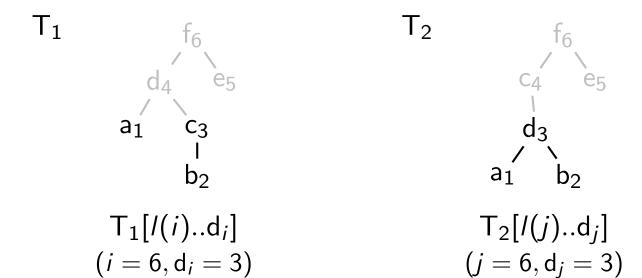


(1) 
$$fdist(T_1[I(i)..d_i - 1], T_2[I(j)..d_j]) + \omega_{del}$$

$$a_1$$
  $c_3$   $d_3$   $b_2$   $b_2$   $T_1[I(i)..d_i - 1]$   $C_1[I(j)..d_j]$ 

- edit script:  $ins(d_3), del(c_3)$
- cost: 1 + 1 = 2

# Example: First Recursive Formula (2/3)



(2) 
$$fdist(T_1[I(i)..d_i], T_2[I(j)..d_j - 1]) + \omega_{ins}$$

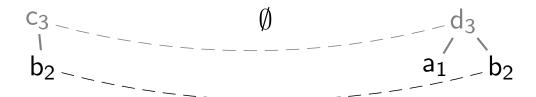
$$a_1$$
  $c_3$   $d_3$   $b_2$   $b_2$   $T_1[I(i)..d_i - 1]$   $C_2[I(j)..d_j]$ 

- edit script:  $del(c_3)$ ,  $ins(d_3)$
- cost: 1 + 1 = 2

# Example: First Recursive Formula (3/3)

(3) 
$$fdist(T_1[I(i)..I(d_i) - 1], T_2[I(j)..I(d_j) - 1]) + fdist(T_1[I(d_i)..d_i - 1], T_2[I(d_j)..d_j - 1]) + \omega_{ren}$$

 $\mathsf{a}_1$ 



$$\mathsf{T}_1[I(i)..I(\mathsf{d}_i)-1] \; \mathsf{T}_1[I(\mathsf{d}_i)..\mathsf{d}_i-1] \; \mathsf{T}_2[I(j)..I(\mathsf{d}_j)-1] \; \mathsf{T}_2[I(\mathsf{d}_j)..\mathsf{d}_j-1]$$

- $T_1[I(i)..d_i 1] \rightarrow T_2[I(j)..d_i 1]$ :  $del(a_1)$
- $T_1[I(d_i)..d_i 1] \rightarrow T_2[I(d_i)..d_i 1]$ :  $ins(a_1)$
- $c_3 \rightarrow d_3$ :  $ren(c_3, d_3)$
- cost: 1+1+1=3

## Analogy to the String Case

• Why is the third formula not (in analogy to the string case):

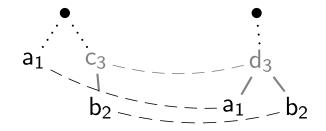
$$fdist(T_1[I(i)..d_i - 1], T_2[I(j)..d_j - 1]) + \omega_{ren}$$

• Consider the previous example:

$$a_1$$
  $c_3$   $a_1$   $b_2$   $a_1$   $b_2$ 

$$T_1[I(i)..d_i - 1] T_2[I(j)..d_i - 1]$$

- $ren(c_3, d_3)$  does not transform  $T_1[I(i)..d_i]$  to  $T_2[I(j)..d_j]$
- In fact the mapping  $M = \{(a_1, a_1), (b_2, b_2), (c_3, d_3)\}$  is not valid:
  - Connect all trees in the forest with a dummy node (●):
  - As  $d_3$  is an ancestor of  $a_1$ ,  $c_3$  must be an ancestor of  $a_1$ , which is false.



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#### Observation

$$fdist(\mathsf{T}_{1}[I(i)..\mathsf{d}_{i}-1],\mathsf{T}_{2}[I(j)..\mathsf{d}_{j}]) + \omega_{del} \\ fdist(\mathsf{T}_{1}[I(i)..\mathsf{d}_{i}],\mathsf{T}_{2}[I(j)..\mathsf{d}_{j}-1]) + \omega_{ins} \\ fdist(\mathsf{T}_{1}[I(i)..I(\mathsf{d}_{i})-1],\mathsf{T}_{2}[I(j)..I(\mathsf{d}_{j})-1]) \\ + fdist(\mathsf{T}_{1}[I(\mathsf{d}_{i})..\mathsf{d}_{i}-1],\mathsf{T}_{2}[I(\mathsf{d}_{j})..\mathsf{d}_{j}-1]) \\ + \omega_{ren} \\$$

- Observation about the First Recursive Formula:
  - $fdist(T_1[I(d_i)..d_i-1], T_2[I(d_i)..d_i-1])$  [D] is a prefix of the subtree rooted in d<sub>i</sub>
  - all other subforests are prefixes of subtrees rooted in i
  - [D] does not fit the scheme (bad for dynamic programming algorithm)
- We derive the Second Recursive Formula:
  - we distinguish two cases (both forests are trees/one forest is not a tree)
  - in each case we replace term [D] by a new term that is easier to handle in a dynamic programming algorithm

#### Second Recursive Formula: Forest Distance

#### Lemma (Second Recursive Formula)

Given two trees  $T_1$  and  $T_2$ ,  $i \in N(T_1)$  and  $d_i \in desc(i)$ ,  $j \in N(T_2)$  and  $d_i \in desc(j)$ , then:

(1) If  $I(i) = I(d_i)$  and  $I(j) = I(d_j)$ , i.e., both forests are trees:

$$fdist(\mathsf{T}_{1}[\mathit{I}(i)..\mathsf{d}_{i}],\mathsf{T}_{2}[\mathit{I}(j)..\mathsf{d}_{j}]) = \min \begin{cases} fdist(\mathsf{T}_{1}[\mathit{I}(i)..\mathsf{d}_{i}-1],\mathsf{T}_{2}[\mathit{I}(j)..\mathsf{d}_{j}]) + \omega_{del} \\ fdist(\mathsf{T}_{1}[\mathit{I}(i)..\mathsf{d}_{i}],\mathsf{T}_{2}[\mathit{I}(j)..\mathsf{d}_{j}-1]) + \omega_{ins} \\ fdist(\mathsf{T}_{1}[\mathit{I}(i)..\mathsf{d}_{i}-1],\mathsf{T}_{2}[\mathit{I}(j)..\mathsf{d}_{j}-1]) + \omega_{ren} \end{cases}$$

(2) If  $I(i) \neq I(d_i)$  and/or  $I(j) \neq I(d_j)$ , i.e., one of the forests is not a tree:

$$fdist(\mathsf{T}_{1}[I(i)..\mathsf{d}_{i}],\mathsf{T}_{2}[I(j)..\mathsf{d}_{j}]) = \min \begin{cases} fdist(\mathsf{T}_{1}[I(i)..\mathsf{d}_{i}-1],\mathsf{T}_{2}[I(j)..\mathsf{d}_{j}]) + \omega_{del} \\ fdist(\mathsf{T}_{1}[I(i)..\mathsf{d}_{i}],\mathsf{T}_{2}[I(j)..\mathsf{d}_{j}-1]) + \omega_{ins} \\ fdist(\mathsf{T}_{1}[I(i)..I(\mathsf{d}_{i})-1],\mathsf{T}_{2}[I(j)..I(\mathsf{d}_{j})-1]) \\ + fdist(\mathsf{T}_{1}[I(\mathsf{d}_{i})..\mathsf{d}_{i}],\mathsf{T}_{2}[I(\mathsf{d}_{j})..\mathsf{d}_{j}]) \end{cases}$$

#### Proof of the Second Recursive Formula

#### Proof.

(1) follows from the previous recursive formula for  $I(i) = I(d_i)$  and  $I(j) = I(d_j)$  as the following holds:

$$fdist(T_1[I(i)..I(d_i) - 1], T_2[I(j)..I(d_i) - 1]) = fdist(\emptyset, \emptyset) = 0.$$

(2) The following inequation holds:

$$[A] \quad fdist(\mathsf{T}_{1}[I(i)..\mathsf{d}_{i}],\mathsf{T}_{2}[I(j)..\mathsf{d}_{j}]) \leq fdist(\mathsf{T}_{1}[I(i)..I(\mathsf{d}_{i})-1],\mathsf{T}_{2}[I(j)..I(\mathsf{d}_{j})-1]) \quad [B] \\ \quad + fdist(\mathsf{T}_{1}[I(\mathsf{d}_{i})..\mathsf{d}_{i}],\mathsf{T}_{2}[I(\mathsf{d}_{j})..\mathsf{d}_{j}]) \qquad [C]$$
 
$$\leq fdist(\mathsf{T}_{1}[I(i)..I(\mathsf{d}_{i})-1],\mathsf{T}_{2}[I(j)..I(\mathsf{d}_{j})-1]) \quad [B] \\ \quad + fdist(\mathsf{T}_{1}[I(\mathsf{d}_{i})..\mathsf{d}_{i}-1],\mathsf{T}_{2}[I(\mathsf{d}_{j})..\mathsf{d}_{j}-1]) \quad [D] \\ \quad + \omega_{ren}$$

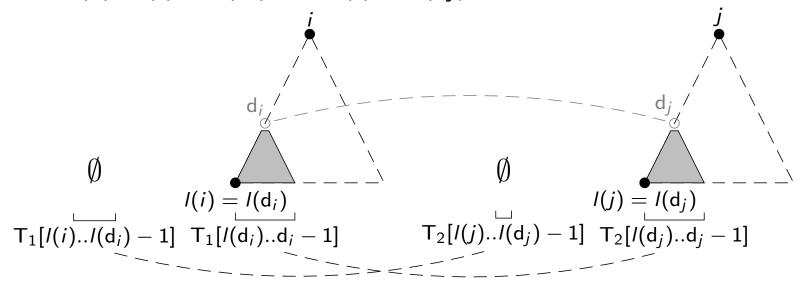
 $A \leq B + C$  as the left-hand side is the *minimal* cost mapping, while the right-hand side is a particular case with a possibly sub-optimal mapping.  $C \leq D + \omega_{ren}$  holds for the same reason.

As we are looking for the *minimum* distance, we can substitute  $D + \omega_{ren}$  by C.



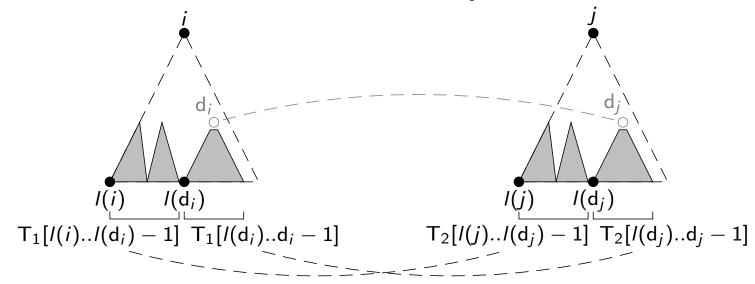
# Illustration: Proof of the Second Recursive Formula (1/2)

• Case (1):  $I(i) = I(d_i)$  and  $I(j) = I(d_j)$ :



# Illustration: Proof of the Second Recursive Formula (2/2)

• Case (2):  $I(i) \neq I(d_i)$  and/or  $I(j) \neq I(d_i)$ :



## Implications by the Second Recursive Formula

• Note:  $fdist(T_1[I(d_i)..d_i], T_2[I(d_j)..d_j]$  is the tree edit distance between the subtrees rooted in  $T[d_i]$  and  $T[d_j]$ . We use the following notation:

$$treedist(d_i, d_j) = fdist(T_1[I(d_i)..d_i], T_2[I(d_j)..d_j])$$

- Dynamic Programming: As the same sub-problem must be solved many times, we use a dynamic programming approach.
- Bottom-Up: As for the computation of the tree distance treedist(i,j) we need almost all values  $treedist(d_i, d_j)$  ( $d_i \in desc(T_1[i])$ ),  $d_j \in desc(T_1[j])$ ), we use a bottom-up approach.
- Key Roots: If
  - $d_i$  is on the path from I(i) to  $T_1[i]$  and
  - $d_j$  is on the path from I(j) to  $T_2[j]$ ,

then  $treedist(d_i, d_j)$  is computed as a byproduct of treedist(i, j). We call the nodes that are *not* computed as a byproducts the key roots.

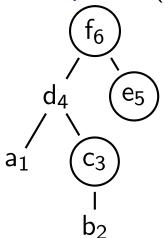
## Key Roots

#### Definition (Key Root)

The set of *key roots* of a tree T is defined as

$$kr(\mathsf{T}) = \{k \in \mathsf{N}(\mathsf{T}) \mid \nexists k' \in \mathsf{N}(\mathsf{T}) : k' > k \text{ and } l(k) = l(k')\}$$

- Alternative definition: A key root is a node of T that either has a left. sibling or is the root of T.
- Example:  $kr(T) = \{3, 5, 6\}$



- Only subtrees rooted in a key root need a separate computation.
- The number of key roots is equal to the number of leaves in the tree.

## Outline

- Tree Edit Distance
  - Preliminaries and Definition
  - Forests Distance and Recursive Formula
  - Second Recursive Formula
  - The Tree Edit Distance Algorithm
  - Example: Tree Edit Distance Computation
  - Complexity of the Tree Edit Distance Algorithm

## The Edit Distance Algorithm I/II

#### tree-edit-dist $(T_1, T_2)$

```
td[1..|\mathsf{T}_1|,1..|\mathsf{T}_2|]: empty array for tree distances; l_1 = \mathsf{Imld}(root(\mathsf{T}_1)); \ kr_1 = \mathsf{kr}(l_1,|leaves(\mathsf{T}_1)|); l_2 = \mathsf{Imld}(root(\mathsf{T}_2)); \ kr_2 = \mathsf{kr}(l_2,|leaves(\mathsf{T}_2)|); for x = 1 to |kr_1| do for y = 1 to |kr_2| do forest-dist(kr_1[x],kr_2[y],l_1,l_2,td);
```

- $l_1$  is an array of size  $|T_1|$ ,  $l_1[i]$  is the leftmost leaf descendant of node i;  $l_2$  is the analog for  $T_2$  (detailed algorithm ImId(.) follows)
- $kr_1$  is an array that contains all the key roots of  $T_1$  sorted in ascending order;  $kr_2$  is the analog for  $T_2$  (detailed algorithm kr(.,.) follows)
- Algorithm and lemmas by [ZS89] (see also [AG97])

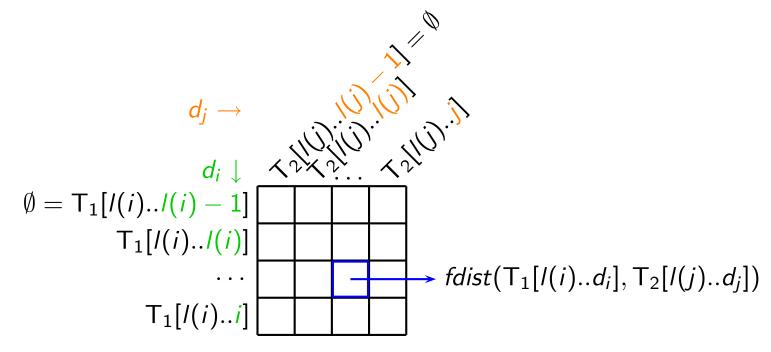
## The Edit Distance Algorithm II/II

#### forest-dist $(i, j, l_1, l_2, td)$

```
fd[l_1[i] - 1..i, l_2[j] - 1..j]: empty array;
fd[l_1[i]-1, l_2[i]-1]=0;
for d_i = l_1[i] to i do fd[d_i, l_2[j] - 1] = fd[d_i - 1, l_2[j] - 1] + \omega_{del};
for d_i = l_2[j] to j do fd[l_1[i] - 1, d_i] = fd[l_1[i] - 1, d_i - 1] + \omega_{ins};
for d_i = l_1[i] to i do
    for d_i = l_2[j] to j do
         if ||f|| = ||f|| and ||f|| = ||f|| then
             fd[d_i, d_i] = \min(fd[d_i - 1, d_i] + \omega_{del},
                                  fd[d_i, d_i - 1] + \omega_{ins}
                                  fd[d_{i}-1,d_{i}-1]+\omega_{ren});
             td[d_i, d_i] = f[d_i, d_i];
         else fd[d_i, d_i] = \min(fd[d_i - 1, d_i] + \omega_{del},
                                     fd[d_i, d_i - 1] + \omega_{ins}
                                     fd[I[d_i] - 1, I[d_i] - 1] + td[d_i, d_i]);
```

### The Temporary Forest Distance Matrix

- $fd[d_i, d_i]$  contains the forest distance between
  - $\mathsf{T}_1[I(i)..d_i]$ , where  $d_i \in desc(\mathsf{T}_1[i])$  and
  - $\mathsf{T}_2[I(j)..d_j]$ , where  $d_j \in desc(\mathsf{T}_2[j])$ .



• fd is temporary and exists only in forest-dist()

### The Tree Distance Matrix

- td[i][j] stores the tree edit distance between
  - the tree rooted in  $T_1[i]$  (i.e.,  $T_1[I(i)..i]$ ) and
  - the tree rooted in  $T_2[j]$  (i.e.,  $T_2[I(j)..j]$ ).
- each call of forest-dist() fills new values into td
- $td[|T_1|, |T_2|]$  stores the tree edit distance between  $T_1$  and  $T_2$

### Computing Key Roots and Left-Most Leaf Descendants

The tree edit distance algorithm uses the following functions:

- ImId(i): computes an array with the left-most leaf descendants of all descendants of a node i
- kr(I, Ic): given the array I = ImId(i) of left-most leaf descendants, and the number Ic of leaf descendants of i, compute all key roots of the subtree rooted in i

#### tree-edit-dist $(T_1, T_2)$

```
 \begin{split} td[1..|\mathsf{T}_1|,1..|\mathsf{T}_2|] : \text{empty array for tree distances;} \\ l_1 &= \mathsf{ImId}(\mathit{root}(\mathsf{T}_1)); \ \mathit{kr}_1 = \mathsf{kr}(\mathit{l}_1,|\mathit{leaves}(\mathsf{T}_1)|); \\ l_2 &= \mathsf{ImId}(\mathit{root}(\mathsf{T}_2)); \ \mathit{kr}_2 = \mathsf{kr}(\mathit{l}_2,|\mathit{leaves}(\mathsf{T}_2)|); \\ \mathbf{for} \ \mathit{x} = 1 \ \mathsf{to} \ |\mathit{kr}_1| \ \mathbf{do} \\ &\quad \mathsf{for} \ \mathit{y} = 1 \ \mathsf{to} \ |\mathit{kr}_2| \ \mathbf{do} \\ &\quad \mathsf{forest-dist}(\mathit{kr}_1[\mathit{x}],\mathit{kr}_2[\mathit{y}],\mathit{l}_1,\mathit{l}_2,\mathit{td}); \end{split}
```

### Computing the Left-Most Leaf Descendants

### ImId(v, I)

```
foreach child c of v (left to right) do I \leftarrow Imld(c, I);
if v is a leaf then
   I[id(v)] \leftarrow id(v)
else
   c_1 \leftarrow \text{ first child of } v;
   I[id(v)] \leftarrow I[id(c_1)];
return I;
```

- Input: root node v of a tree T, empty array /[1..|T|]
- Output: array I, I[i] is the left-most leaf descendent of node T[i]

### Computing the Key Roots

### kr(I, Ic)

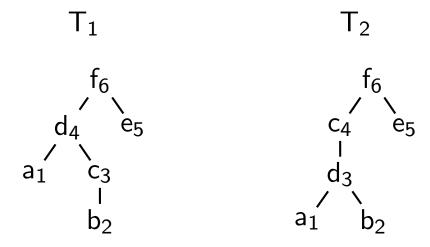
```
kr[1..lc]: empty array;
visited[]: boolean array of size |/|, init with false;
k \leftarrow |kr|; \quad i \leftarrow |I|;
while k > 1 do
        if not visited[/[i]] then
           kr[k--] \leftarrow i;
           visited[I[i]] \leftarrow true;
       ī−-;
return sort(kr);
```

- Input:
  - /[1..|T|]: /[i] is the left-most leaf descendent of node T[i]
  - lc = |leaves(T)| is the number of leaves in T
- Output: array kr[1..|leaves(T)|] with key roots sorted by node ID
- Note: Loop condition is correct, as  $k \ge 1 \Rightarrow i \ge 1$ (the number of key roots is exactly the number of leaves, and it will always be filled when all nodes are traversed)

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## Example Trees and Edit Costs



- Example: Edit distance between  $T_1$  and  $T_2$ .
  - ullet  $\omega_{\mathit{ins}} = \omega_{\mathit{del}} = 1$
  - $\bullet$   $\omega_{ren}=0$  for identical rename, otherwise  $\omega_{ren}=1$
- Each of the following slide is the result of a call of forest-dist().

# Executing the Algorithm (1/9)

• 
$$i = kr_1[x] = 3 \Rightarrow l_1[i] = 2$$

• 
$$j = kr_2[y] = 2 \Rightarrow l_2[j] = 2$$

temporary array fd:

$$\begin{array}{c|cccc} \mathsf{d}_j & \longrightarrow & 2 \\ \mathsf{d}_i \downarrow & \boxed{0} & \boxed{1} \\ 2 & \boxed{1} & \boxed{0} \\ 3 & \boxed{2} & \boxed{1} \end{array}$$

$$I_1[i] = I_1[d_i]$$
 and  $I_2[j] = I_2[d_i]$ 

 $I_2$  $kr_2$ 

	1	2	3	4	5	6
1						
2		0				
3		1				
2 3 4 5 6						
5						
6						

# Executing the Algorithm (2/9)

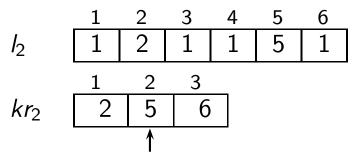
• 
$$i = kr_1[x] = 3 \Rightarrow l_1[i] = 2$$

• 
$$j = kr_2[y] = 5 \Rightarrow l_2[j] = 5$$

temporary array fd:

$$\begin{array}{c|cccc} \mathsf{d}_j \to & 5 \\ \mathsf{d}_i \downarrow & 0 & 1 \\ 2 & 1 & 1 \\ 3 & 2 & 2 \end{array}$$

$$I_1[i] = I_1[d_i]$$
 and  $I_2[j] = I_2[d_i]$ 



	1	2	3	4	5	6
1						
2		0			1	
<ul><li>2</li><li>3</li><li>4</li><li>5</li><li>6</li></ul>		1			2	
4						
5						
6						

# Executing the Algorithm (3/9)

• 
$$i = kr_1[x] = 3 \Rightarrow l_1[i] = 2$$

• 
$$j = kr_2[y] = 6 \Rightarrow l_2[j] = 1$$

• temporary array *fd*:

(	$d_j  o$	1	2	3	4	5	6
$d_i \downarrow$	0	1	2	3	4	5	6
2	1	1	1	2	3	4	5
3	2	2	2	2	2	3	4

$$I_1[i] = I_1[d_i]$$
 and  $I_2[j] = I_2[d_j]$ 

_				
	1	2	3	
$kr_2$	2	5	6	
			<u> </u>	

	1	2	3	4	5	6
1						
2 3	1	0	2	3	1	5
3	2	1	2	2	2	4
4						
5						
6						

# Executing the Algorithm (4/9)

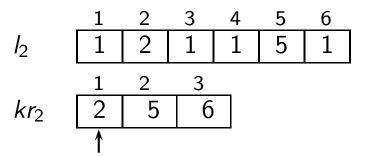
• 
$$i = kr_1[x] = 5 \Rightarrow l_1[i] = 5$$

• 
$$j = kr_2[y] = 2 \Rightarrow l_2[j] = 2$$

• temporary array fd:

$$\begin{array}{c|cccc}
d_j & \rightarrow & 2 \\
d_i \downarrow & 0 & 1 \\
5 & 1 & 1
\end{array}$$

$$I_1[i] = I_1[d_i]$$
 and  $I_2[j] = I_2[d_j]$ 



	1	2	3	4	5	6
1						
2 3	1	0	2	3	1	5
3	2	1	2	2	2	4
4						
5		1				
6		·	·			

# Executing the Algorithm (5/9)

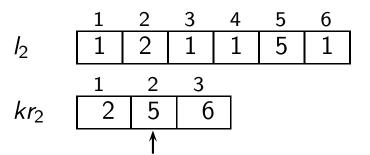
• 
$$i = kr_1[x] = 5 \Rightarrow l_1[i] = 5$$

• 
$$j = kr_2[y] = 5 \Rightarrow l_2[j] = 5$$

• temporary array fd:

$$d_i \downarrow \begin{array}{c|c} d_j \rightarrow & 5 \\ \hline 0 & 1 \\ \hline 5 & 1 & 0 \\ \hline \end{array}$$

$$I_1[i] = I_1[d_i] \text{ and } I_2[j] = I_2[d_j]$$



	1	2	3	4	5	6
1						
2 3	1	0	2	3	1	5
3	2	1	2	2	2	4
4						
5 6		1			0	
6						

# Executing the Algorithm (6/9)

• 
$$i = kr_1[x] = 5 \Rightarrow l_1[i] = 5$$

• 
$$j = kr_2[y] = 6 \Rightarrow l_2[j] = 1$$

• temporary array fd:

	J				4		
d <sub>i</sub> ↓ 5	0	1	2	3	4	5	6
5	1	1	2	3	4	4	5

$$I_1[i] = I_1[d_i]$$
 and  $I_2[j] = I_2[d_i]$ 

		2	3	4	5	O
$I_2$	1	2	1	1	5	1
	1	2	3			
kr <sub>2</sub>	2	5	6			
			1			

	1	2	3	4	5	6
1						
2 3	1	0	2	3	1	5
3	2	1	2	2	2	4
4						
5	1	1	3	4	0	5
6						

# Executing the Algorithm (7/9)

• 
$$i = kr_1[x] = 6 \Rightarrow l_1[i] = 1$$

• 
$$j = kr_2[y] = 2 \Rightarrow l_2[j] = 2$$

• temporary array *fd*:

$$I_1[i] = I_1[d_i] \text{ and } I_2[j] = I_2[d_j]$$

	1	2	3	4	5	6
1		1				
2 3	1	0	2	3	1	5
3	2	1	2	2	2	4
4		3				
5	1	1	3	4	0	5
6		5				

# Executing the Algorithm (8/9)

• 
$$i = kr_1[x] = 6 \Rightarrow l_1[i] = 1$$

• 
$$j = kr_2[y] = 5 \Rightarrow l_2[j] = 5$$

• temporary array *fd*:

$$I_1[i] = I_1[d_i] \text{ and } I_2[j] = I_2[d_j]$$

	1	2	3	4	5	6
1		1			1	
2 3	1	0	2	3	1	5
3	2	1	2	2	2	4
4		3			4	
5	1	1	3	4	0	5
6		5			5	

# Executing the Algorithm (9/9)

$$l_2$$
 $1 \quad 2 \quad 1 \quad 1 \quad 5 \quad 1$ 
 $kr_2$ 
 $2 \quad 5 \quad 6$ 

• 
$$i = kr_1[x] = 6 \Rightarrow l_1[i] = 1$$

• 
$$j = kr_2[y] = 6 \Rightarrow l_2[j] = 1$$

• temporary array *fd*:

(	$d_j  o$	1	2	3	4	5	6
$d_i \downarrow$	0	1	2	3	4	5	6
1	1	0	1	2	3	4	5
2	2	1	0	1	2	3	4
3	3	2	1	2	3	4	5
4	4	3	2	1	2	3	4
5	5	4	3	2	3	2	3
6	6	5	4	3	3	3	2

$$I_1[i] = I_1[d_i] \text{ and } I_2[j] = I_2[d_j]$$

	1	2	3	4	5	6
1	0	1	2	3	1	5
2 3	1	0	2	3	1	5
3	2	1	2	2	2	4
4	3	3	1	2	4	4
5	1	1	3	4	0	5
6	5	5	3	3	5	2

### Outline

- Tree Edit Distance
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### Notation

- Notation:
  - |T| is the number of nodes in T
  - depth(v) is the number of ancestors of v (including v itself)
  - depth(T) is the maximum depth of a node in T
  - leaves(T) is the set of leaves of T
  - t(i) is the subtree rooted in node i

### forest-dist: Time Complexity

### forest-dist $(i, j, l_1, l_2, td)$

```
 \begin{split} fd[l_1[i]-1..i,l_2[j]-1..j] : & \text{empty array}; \\ fd[l_1[i]-1,l_2[j]-1] = 0; \\ \textbf{for } d_i = l_1[i] \textbf{ to } i \textbf{ do } fd[d_i,l_2[j]-1] = fd[d_i-1,l_2[j]-1] + \omega_{del}; \\ \textbf{for } d_j = l_2[j] \textbf{ to } j \textbf{ do } fd[l_1[i]-1,d_j] = fd[l_1[i]-1,d_j-1] + \omega_{ins}; \\ \textbf{for } d_i = l_1[i] \textbf{ to } i \textbf{ do } \\ \textbf{ for } d_j = l_2[j] \textbf{ to } j \textbf{ do } \\ \textbf{ if } l[d_i] = l[i] \textbf{ and } l[d_j] = l[j] \textbf{ then } \\ fd[d_i,d_j] = \min(\ldots); \\ td[d_i,d_j] = f[d_i,d_j]; \\ \textbf{ else } fd[d_i,d_j] = \min(\ldots); \end{split}
```

- Input nodes are i and j.
- They are root nodes of subtrees  $t_1(i)$  and  $t_2(j)$ .
- The nested loop is executed  $|t_1(i)| \times |t_2(j)|$  times.
- $\Rightarrow$  Time complexity  $O(|t_1(i)| \times |t_2(j)|)$

### tree-edit-dist: Time Complexity

#### tree-edit-dist $(T_1, T_2)$

```
td[1..|\mathsf{T}_1|,1..|\mathsf{T}_2|]: empty array for tree distances; l_1 = \mathsf{Imld}(root(\mathsf{T}_1)); \ kr_1 = \mathsf{kr}(l_1,|leaves(\mathsf{T}_1)|); l_2 = \mathsf{Imld}(root(\mathsf{T}_2)); \ kr_2 = \mathsf{kr}(l_2,|leaves(\mathsf{T}_2)|); for x = 1 to |kr_1| do forest-dist(kr_1[x],kr_2[y],l_1,l_2,td);
```

- Computing  $I_{1/2}$  and  $kr_{1/2}$  is linear,  $O(|\mathsf{T}_1| + |\mathsf{T}_2|)$
- Main loop executes forest-dist()  $|kr_1| \times |kr_2|$  times.
- Complexity:

$$\sum_{i \in kr_1} \sum_{j \in kr_2} |t_1(i)| \times |t_2(j)| = \sum_{i \in kr_1} |t_1(i)| \times \sum_{j \in kr_2} |t_2(j)|$$

• The following lemmas help us to reformulate this expression.

### Collapsed Depth

• **Definition:** The collapsed depth of a node v in T is

$$cdepth(v) = |anc(v) \cap kr(T)|,$$

i.e., the number of ancestors of v (including v itself) that are key roots.

## Collapsed Depth

#### Lemma (Collapsed Depth)

For a tree T with key roots kr(T)

$$\sum_{k \in kr(\mathsf{T})} |t(k)| = \sum_{k=1}^{|\mathsf{T}|} cdepth(k)$$

#### Proof.

- Consider the left-hand formula:
  - A node i of T is counted whenever it appears in a subtree t(k).
  - Node i is in the subtree t(k) iff k is the ancestor of i.
  - Only the subtrees of key roots are considered.
- Thus a node *i* is counted once for each ancestor key root.
- cdepth(i) is the number of ancestor key roots of i (definition of collapsed depth).



### Collapsed Depth

Now we can rewrite the complexity formula:

$$\sum_{i \in kr_1} |t_1(i)| \times \sum_{j \in kr_2} |t_2(j)| = \sum_{i=1}^{|\mathsf{T}_1|} cdepth(i) \times \sum_{j=1}^{|\mathsf{T}_2|} cdepth(j)$$

•  $cdepth(T) \ge cdepth(k)$  for a node k of T, thus

$$\sum_{i=1}^{|\mathsf{T}_1|} cdepth(i) \times \sum_{j=1}^{|\mathsf{T}_2|} cdepth(j) \leq |\mathsf{T}_1| |\mathsf{T}_2| cdepth(\mathsf{T}_1) cdepth(\mathsf{T}_2)$$

- Two obvious upper bounds for the collapsed depth:
  - the tree depth:  $cdepth(T) \leq depth(T)$
  - the number of key roots:  $cdepth(T) \leq |kr(T)|$
- We show that the number of key roots matches the number of leaves.

### Number of Key Roots

#### Lemma (Number of Key Roots)

The number of key roots of a tree is equal to the number of leaves:

$$|kr(\mathsf{T})| = |leaves(\mathsf{T})|$$

#### Proof.

We show that I() is a bijection from the key roots kr(T) to the leaves(T):

- (a) **Injection** for any  $i, j \in kr(T)$ ,  $i \neq j \Rightarrow l(i) \neq l(j)$ : If i > j and l(i) = l(j), j can not be a key root by definition. Analogous rational hold for j > i.
- (b) **Surjection** Each leaf x has a key root  $i \in kr(T)$  such that I(i) = x: If there is no node i > x with I(i) = I(x), then by definition x itself is a key root (I(x) = x) is always true. Otherwise i is the key root of x.



### Complexity of the Tree Edit Distance Algorithm

#### Theorem (Complexity of the Tree Edit Distance Algorithm)

Let  $D_1$  and  $D_2$  denote the depth,  $L_1$  and  $L_2$  the number of leave nodes, and  $N_1$  and  $N_2$  the total number of nodes of two trees  $T_1$  and  $T_2$ , respectively.

(1) The runtime of the tree edit distance algorithm is

$$O(N_1N_2 \min(D_1, L_1) \min(D_2, L_2)).$$

- (2) Let  $N = \max(N_1, N_2)$ . For full, balanced, binary trees the runtime is  $O(N^2 \log^2 N)$ .
- (3) In the worst case min(D, L) = O(N) and the runtime is  $O(N^4)$ .
- (4) The algorithm needs  $O(N_1N_2)$  space.

## Proof of the Complexity Theorem

#### Proof.

- (1) Runtime (general formula): We have shown before, that the complexity is  $O(|T_1||T_2|cdepth(T_1) cdepth(T_2))$ . As  $cdepth(T) \leq |kr(T)| = |leaves(T)|$  (see definition of cdepth(T) and previous lemma) and  $cdepth(T) \leq depth(T)$  (follows from the definition of cdepth(T)), if follows that  $cdepth(T) \leq min(depth(T), |leaves(T)|)$ .
- (2) Full, balanced, binary trees: In this case  $depth(T) = O(\log(|T|))$ .
- (3) Worst case: A full binary tree (i.e., each node has zero or two children) where each non-leaf nodes has at least one leaf child:  $\min(depth(T), |leaves(T)|) = O(|T|)$ .
- (4) Space: The size of the tree distance matrix td is  $|T_1| \times |T_2|$ . In each call of forest-dist() we need a matrix of size  $O(|T_1| \times |T_2|)$ , which is freed when we exit the subroutine.

### Recent Improvements of the Complexity

- Klein [Kle98] improves the worst case for the runtime to  $O(|T_1|^2|T_2|\log(|T_2|))$ , thus from  $O(N^4)$  to  $O(N^3\log(N))$ .
- Dulucq and Touzet [DT03] also give an  $O(N^3 \log(N))$  algorithm.
- Demaine et al. [DMRW07] give an  $O(N^3)$  algorithm. They show that the algorithm is optimal among decomposition algorithms (algorithms as in [ZS89, Kle98, DT03]), i.e., the lower bound is also  $\Omega(N^3)$ .

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