



Department of Computer Science

# Near-optimal labeling schemes for nearest common ancestors

Stephen Alstrup\*

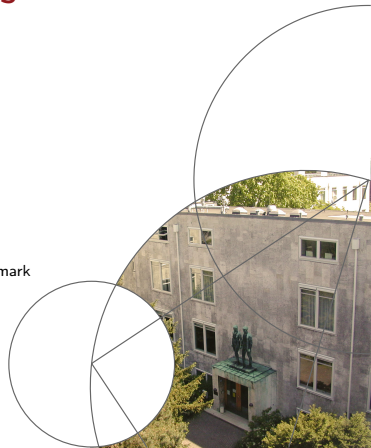
*Esben Bistrup Halvorsen\**

Kasper Green Larsen†

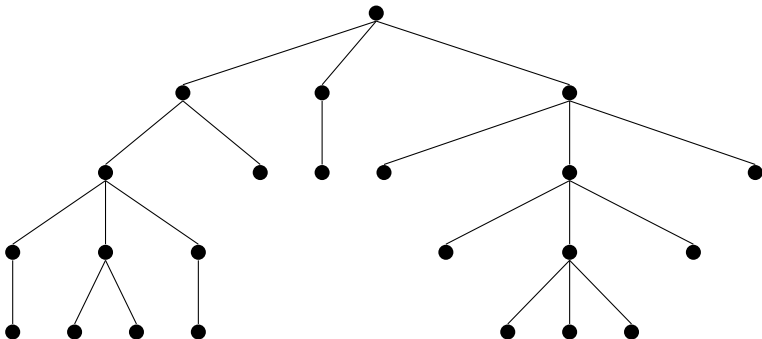
\* Department of Computer Science, University of Copenhagen, Denmark

† MADALGO, Department of Computer Science, Aarhus University, Denmark

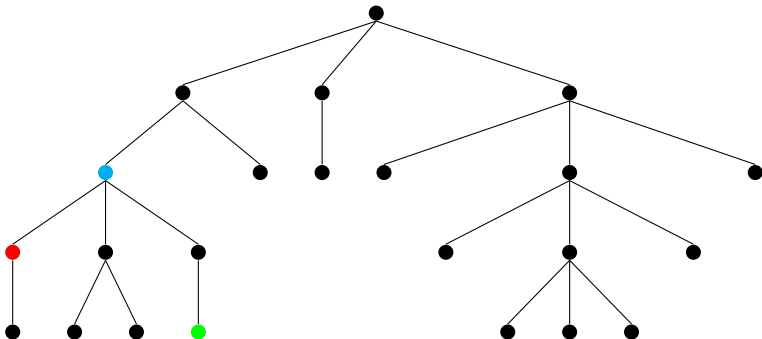
January 6, 2014



# Nearest common ancestor (NCA)



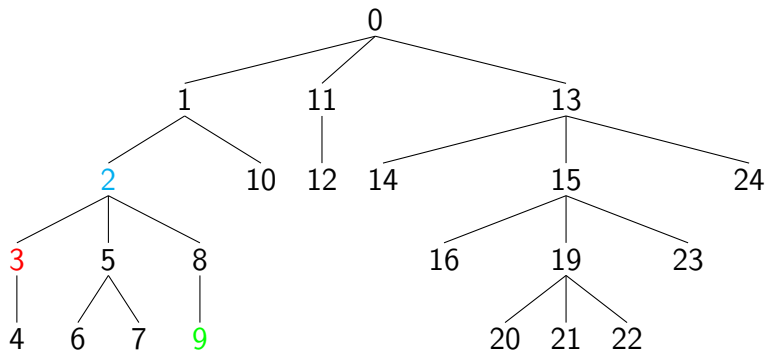
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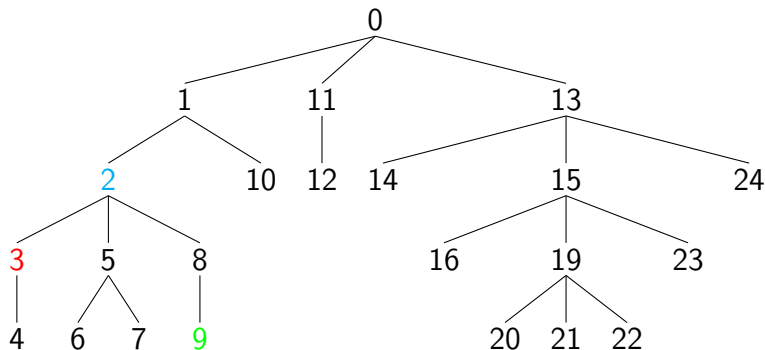
$$\text{nca}(\text{red}, \text{green}) = \text{blue}$$



# NCA labeling schemes



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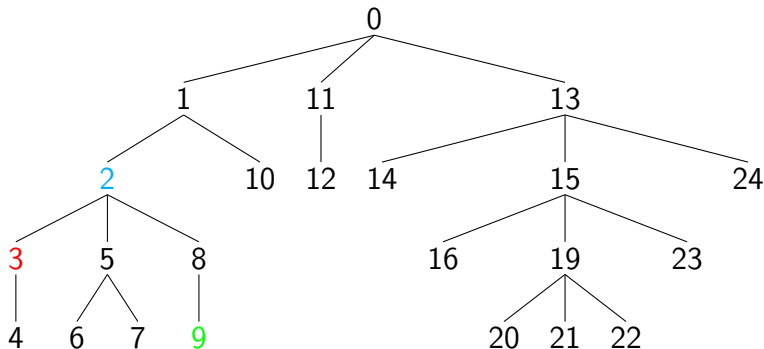
$$l(\bullet) = (0, 1, 2, 3)$$

$$l(\bullet) = (0, 1, 2, 8, 9)$$

$$l(\bullet) = (0, 1, 2)$$



# NCA labeling schemes



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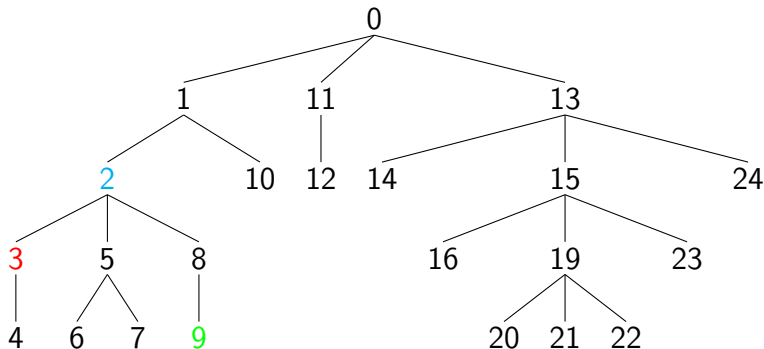
$$l(\bullet) = (0, 1, 2, 8, 9)$$

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$$\max |l(v)| = O(n \log n)$$



# NCA labeling schemes



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**Objective:** minimize  $\max |l(v)|$



# Previous results

## Algorithms to compute NCAs





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- different settings (static, dynamic, . . . )



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- different applications in mind (computational biology, XML, routing, . . . )



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\* see survey by Alstrup, Gavaille, Kaplan, Rauhe (TOCS, 2004)



# Previous results

## Labeling schemes



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- adjacency, ancestry, distance, routing, NCA, ...



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- adjacency, ancestry, distance, routing, NCA, ...
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## Labeling schemes\*

- adjacency, ancestry, distance, routing, NCA, ...
- static/dynamic
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\* see survey by Gavoille, Peleg (Dist. Comp., 2003)



# Previous results

**Harel, Tarjan (Siam J. Computing, 1984):**

NCAs in constant time with linear preprocessing time.

(Includes a labeling scheme for complete binary trees.)



# Previous results

**Kannan, Naor, Rudich (STOC, 1988):**  
Adjacency and ancestry labeling schemes.



# Previous results

**Alstrup, Gavoille, Kaplan, Rauhe (TOCS, 2004):**  
NCA labeling scheme with label size  $10 \log n$ .



# Previous work

**Fischer (ESA, 2009):**

Experimental results with label size  $8 \log n$ .



# Previous results

**Alstrup, Bille, Rauhe (Siam J. Discrete Math., 2005):**  
NCA labeling schemes have label size  $\log n + \Omega(\log \log n)$ .



# Previous results

**Fraigniaud, Korman (STOC, 2010):**

Ancestor labeling schemes of  $\log n + \Theta(\log \log n)$





# New results

## **Alstrup, Halvorsen, Larsen (SODA, 2014):**

Upper bound of  $2.772 \log n$  (and  $2.585 \log n$  for binary trees).

Efficient upper bound of  $3 \log n$ .

Lower bound of  $1.008 \log n$ .



# New results

Labels have size  $(2 \pm \epsilon) \log n$ ,  $\epsilon < 1$ .



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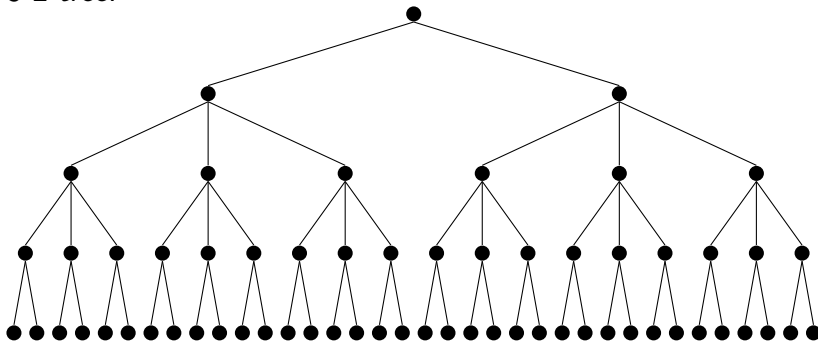
- Separates NCA from ancestry
- First lower bound of  $\log n + \omega(\log \log n)$  for non-distance labeling schemes for trees
- More efficient than previous experimental results
- Total space comparable to “normal” NCA algorithms



3-2 sequence:

$$(2, 3, 3, 2)$$

3-2 tree:

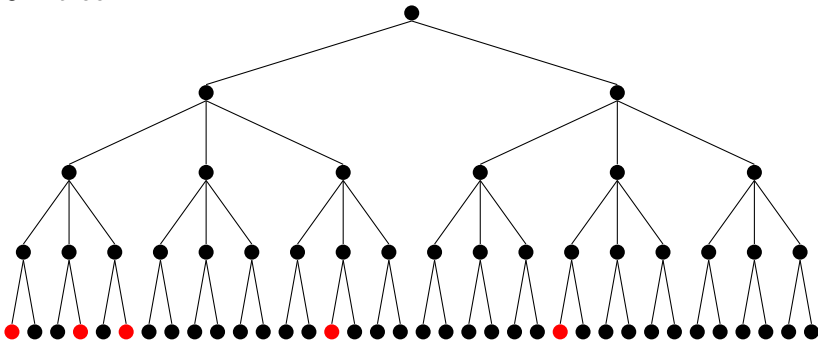


# Lower bound technique

3-2 sequence:

$(2, 3, 3, 2)$

3-2 tree:



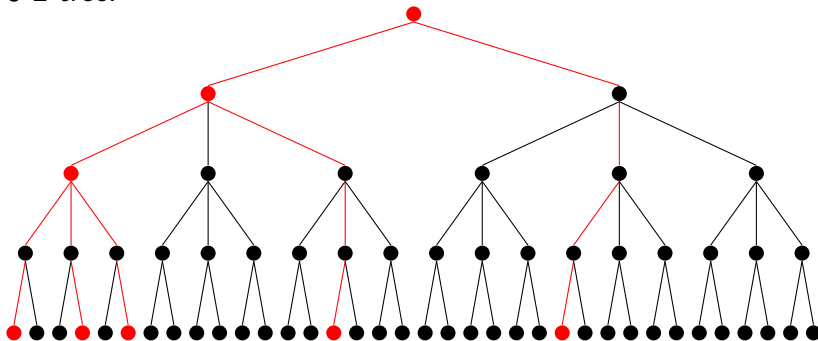


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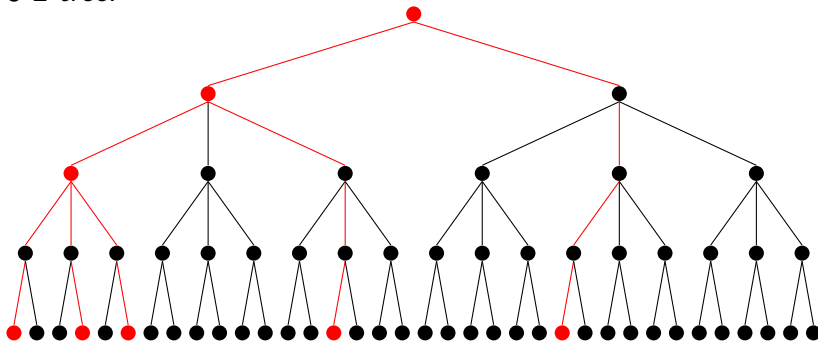
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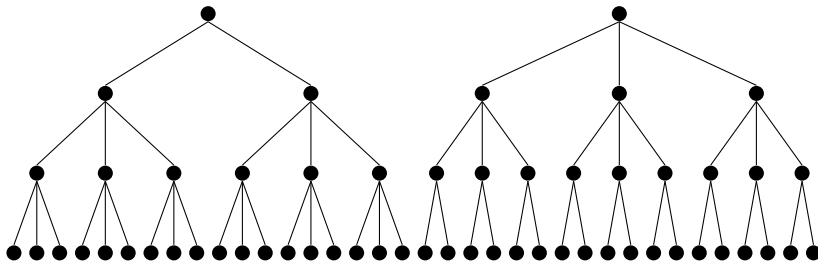
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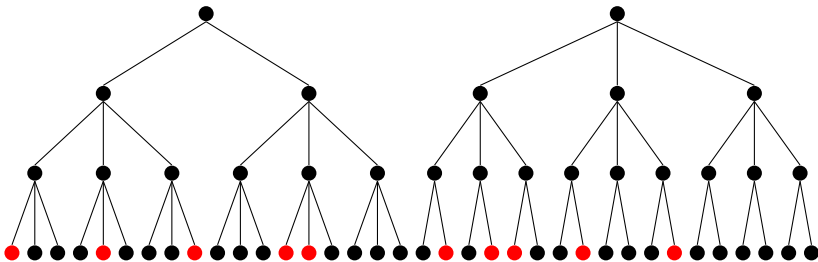
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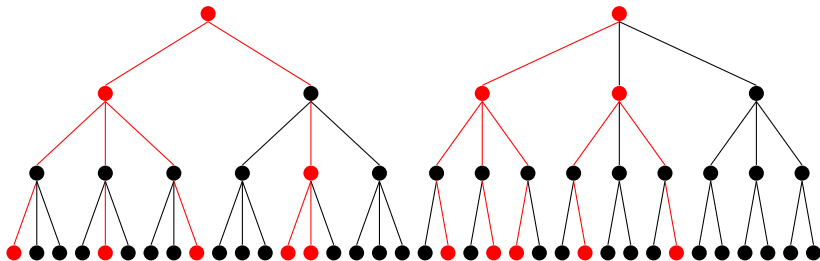
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# Lower bound technique

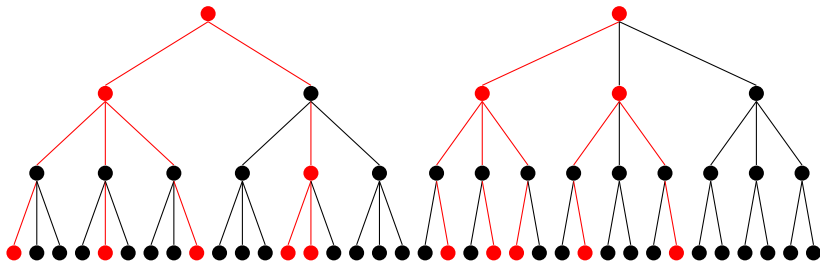
3-2 sequences:

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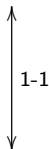
$(3, 3, 2)$

3-2 trees:



# Lower bound technique

$\{(3-2)\text{-sequences with pairwise distance} > h\}$

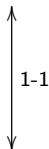


$\{(3-2)\text{-trees with pairwise fewer than } m \text{ labels in common}\}$



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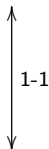
$\Rightarrow [\dots, \textit{math}, \textit{math}, \textit{math}, \dots]$





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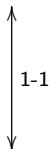
$\Rightarrow [\dots, \textit{math}, \textit{math}, \textit{math}, \dots]$

$\Rightarrow |\{\text{labels}\}| \geq n^{1.008}$



# Lower bound technique

$\{(3-2)\text{-sequences with pairwise distance} > h\}$



$\{(3-2)\text{-trees with pairwise fewer than } m \text{ labels in common}\}$

$\Rightarrow [\dots, \textit{math}, \textit{math}, \textit{math}, \dots]$

$\Rightarrow |\{\text{labels}\}| \geq n^{1.008}$

$\Rightarrow$  labels have worst-case size  $1.008 \log n$



# Future studies



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- Tighter bounds than  $> 1.008 \log n$  and  $< 2.772 \log n$ .



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- Time-efficient version



# Future studies

- Tighter bounds than  $> 1.008 \log n$  and  $< 2.772 \log n$ .
- Time-efficient version
- Dynamic, probabilistic, ...



# Thanks!

