

# ALGORITHMS OF BIOINFORMATICS

# 6

## Suffix Trees

*11 December 2025*

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## Outline

# 6 Suffix Trees

- 6.1 Suffix Trees
- 6.2 Applications
- 6.3 Longest Common Extensions
- 6.4 Suffix Arrays
- 6.5 Suffix sorting: Induced sorting and merging
- 6.6 Suffix Sorting: The DC3 Algorithm
- 6.7 The LCP Array
- 6.8 LCP Array Construction

# Context

*We're still working towards practical solutions for the read mapping problem.*

*So far, our preprocessing was mostly getting smart on the **reads/patterns**.*

- ~~ Now preprocess the genome/text.

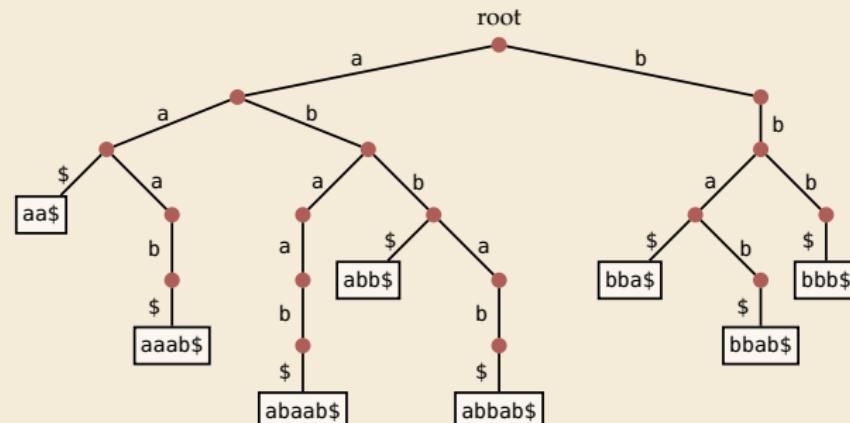
## 6.1 Suffix Trees

# Recap: Tries

- efficient dictionary data structure for strings (or for Aho-Corasick automata 😊)
- name from retrieval, but pronounced “try”
- tree based on symbol comparisons
- **Assumption here:** stored strings are *prefix-free* (no string is a prefix of another)
  - strings of same length ✓
  - some character  $\notin \Sigma$
  - strings have “end-of-string” marker \$ ✓

## ► Example:

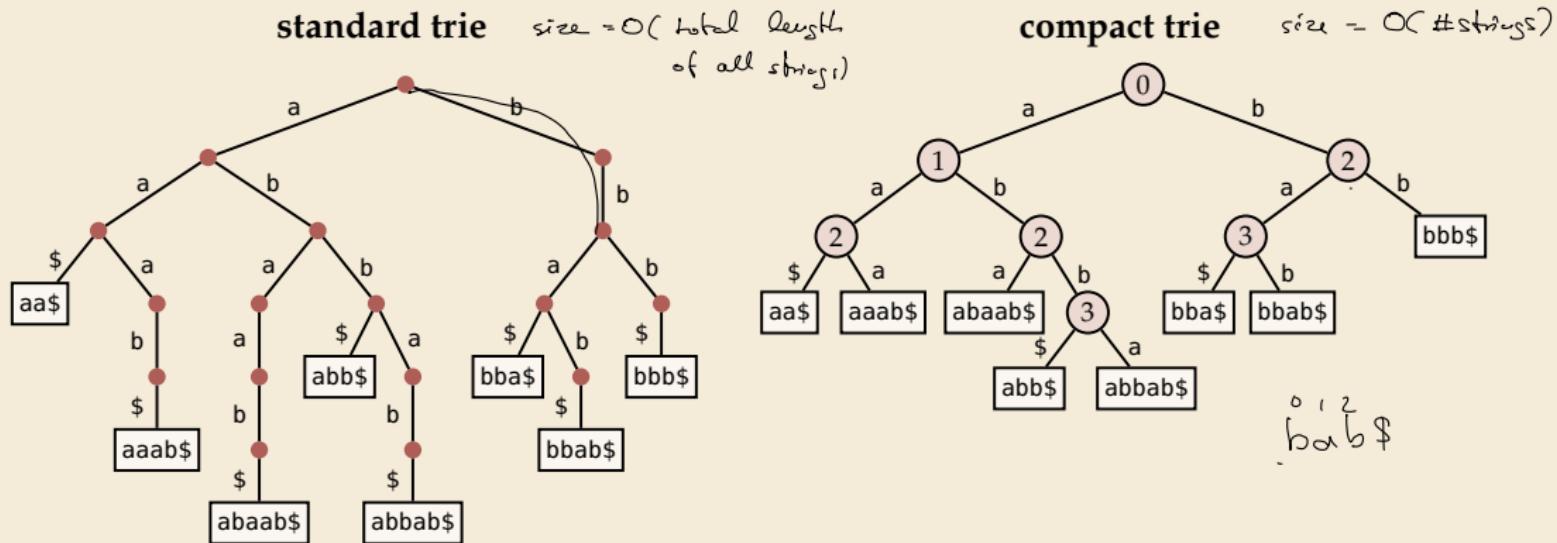
{aa\$, aaab\$, abaab\$, abb\$,  
abbab\$, bba\$, bbab\$, bbb\$}



# Compact tries

- ▶ compress paths of unary nodes into single edge
- ▶ nodes store *index* of next character to check

=1 child



- ▶ search gives first character of edge only ↳ must check for match against stored string
- ▶ all nodes  $\geq 2$  children ↳  $\#nodes \leq \#leaves = \#strings$  ↳ linear space

# Suffix trees – A ‘magic’ data structure

**Appetizer:** Longest common substring problem

- ▶ Given: strings  $S_1, \dots, S_k$                            **Example:**  $S_1 = \text{superiorcalifornialives}$ ,  $S_2 = \text{sealiver}$
- ▶ Goal: find the longest substring that occurs in all  $k$  strings

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Enter: *suffix trees*

- ▶ versatile data structure for index with full-text search
- ▶ linear time (for construction) and linear space
- ▶ allows efficient solutions for many advanced string problems



*“Although the longest common substring problem looks trivial now, given our knowledge of suffix trees, it is very interesting to note that in 1970 Don Knuth conjectured that a linear-time algorithm for this problem would be impossible.”*

[Gusfield: *Algorithms on Strings, Trees, and Sequences* (1997)]

## Suffix trees – Definition

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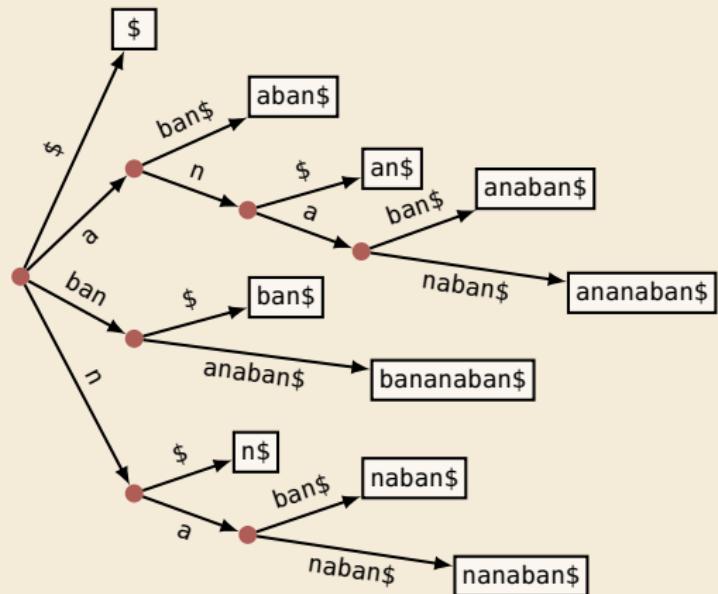
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Example:

$T = \text{bananaban\$}$

suffixes:  $\{\text{bananaban\$}, \text{ananaban\$}, \text{nanaban\$}, \text{anaban\$}, \text{naban\$}, \text{aban\$}, \text{ban\$}, \text{an\$}, \text{n\$}, \$\}$

$T = \begin{array}{cccccccccc} 0 & 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 & 9 \\ \boxed{\text{b}} & \boxed{\text{a}} & \boxed{\text{n}} & \boxed{\text{a}} & \boxed{\text{n}} & \boxed{\text{a}} & \boxed{\text{b}} & \boxed{\text{a}} & \boxed{\text{n}} & \boxed{\$} \end{array}$



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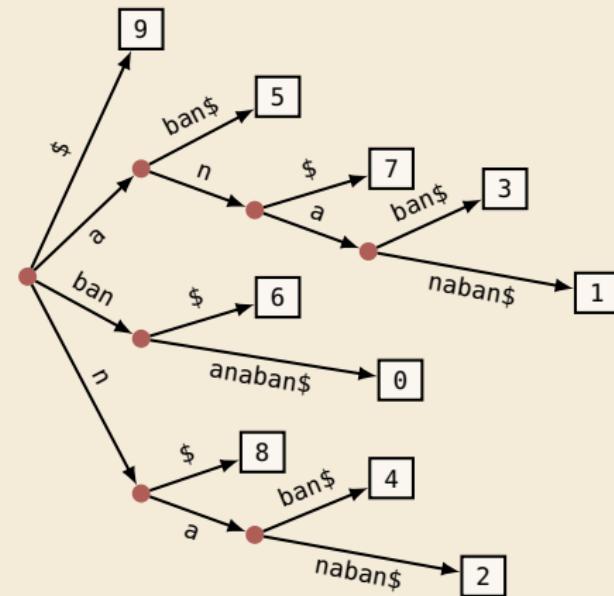
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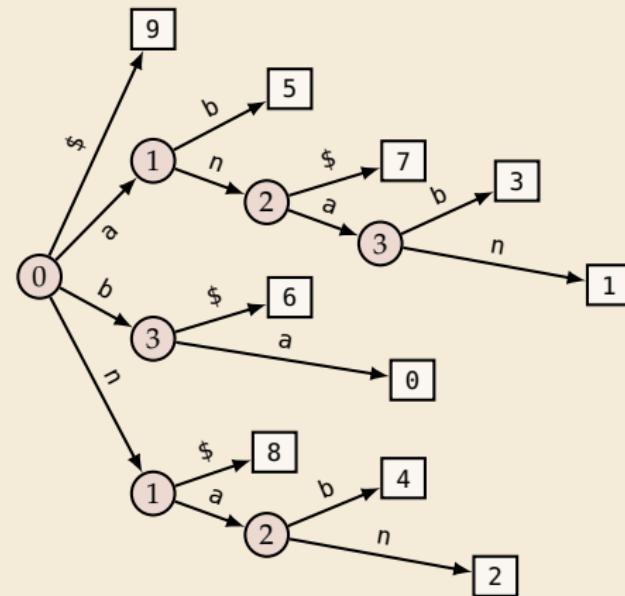
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- also: edge labels like in compact trie
- (more readable form on slides to explain algorithms)



## Suffix trees – Construction

- ▶  $T[0..n]$  has  $n + 1$  suffixes (starting at character  $i \in [0..n]$ )
- ▶ We can build the suffix tree by inserting each suffix of  $T$  into a compressed trie.  
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same order of growth as reading the text!  
Amazing result: Can construct the suffix tree of  $T$  in  $\Theta(n)$  time!

- ▶ several fundamentally different methods known
- ▶ started as theoretical breakthrough
- ▶ now routinely used in bioinformatics practice

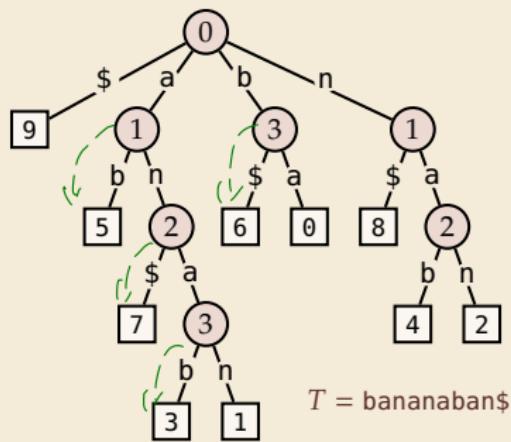
↷ for now, take linear-time construction for granted. What can we do with them?

## 6.2 Applications

# Applications of suffix trees

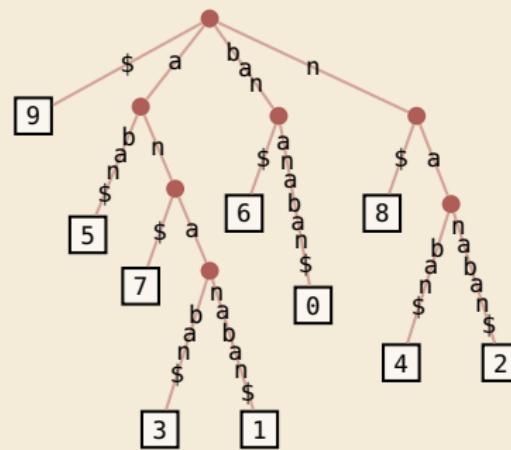
- In this section, always assume suffix tree  $\mathcal{T}$  for  $T$  given.

Recall:  $\mathcal{T}$  stored like this:



$$T = \text{bananaban\$}$$

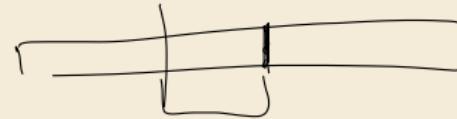
but think about this:



- Moreover: assume internal nodes store pointer to *leftmost leaf in subtree*.
- Notation:  $T_i = T[i..n]$  (including \$)

## Application 1: Text Indexing / String Matching

- ▶  $P$  occurs in  $T \iff P$  is a prefix of a suffix of  $T$
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~~ (try to) follow path with label  $P$ , until

1. we get stuck

at internal node (no node with next character of  $P$ )  
or inside edge (mismatch of next characters)

~~  $P$  does not occur in  $T$

2. we run out of pattern

reach end of  $P$  at internal node  $v$  or inside edge towards  $v$   
~~  $P$  occurs at all leaves in subtree of  $v$

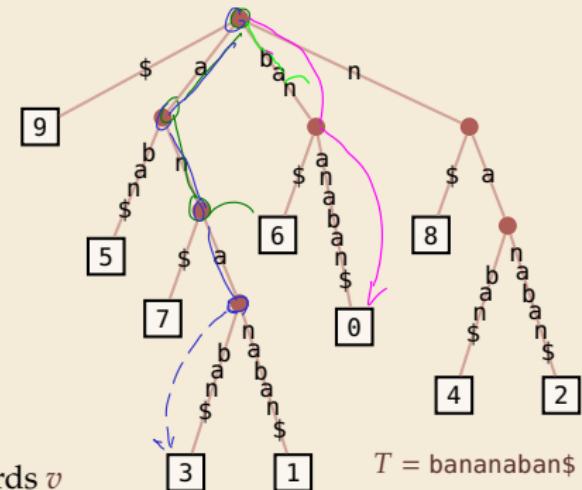
3. we run out of tree

reach a leaf  $\ell$  with part of  $P$  left ~~ compare  $P$  to  $\ell$ .



This cannot happen when testing edge labels since  $\$ \notin \Sigma$ ,  
but needs check(s) in compact trie implementation!

► Finding first match (or NO\_MATCH) takes  $O(|P|)$  time!



Examples:

►  $P = \underline{\text{ann}}$

►  $P = \underline{\text{baa}}$

►  $P = \underline{\text{ana}}$

►  $P = \underline{\text{ba}}$

►  $P = \underline{\text{briar}}$

## Application 2: Longest repeated substring

► **Goal:** Find longest substring  $T[i..i + \ell]$  that occurs also at  $j \neq i$ :  $T[j..j + \ell] = T[i..i + \ell]$ .



~~e.g. for compression  $\rightsquigarrow$  Unit 7~~

? How can we efficiently check *all possible substrings*?

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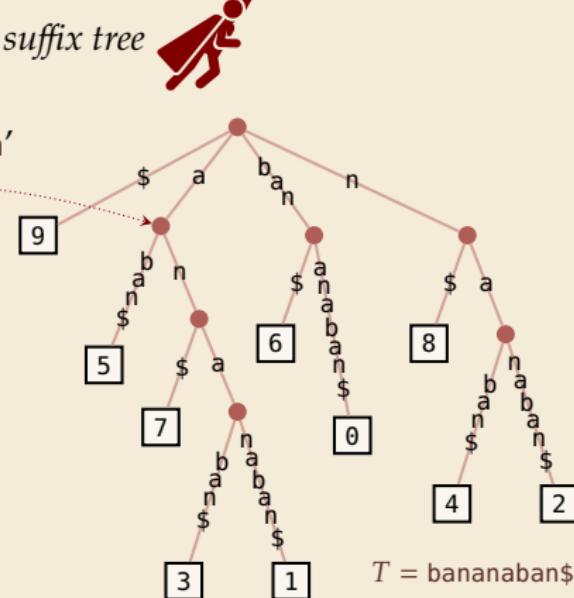
Repeated substrings = shared paths in *suffix tree*



- $T_5 = \text{aban\$}$  and  $T_7 = \text{an\$}$  have *longest common prefix* 'a'

$\rightsquigarrow \exists$  internal node with path label 'a'

here single edge, can be longer path



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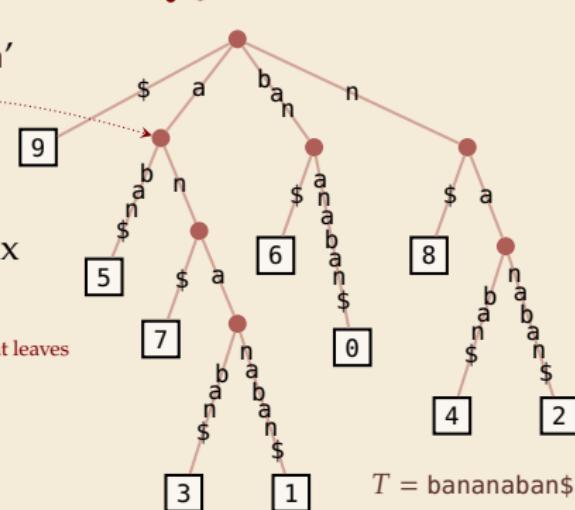
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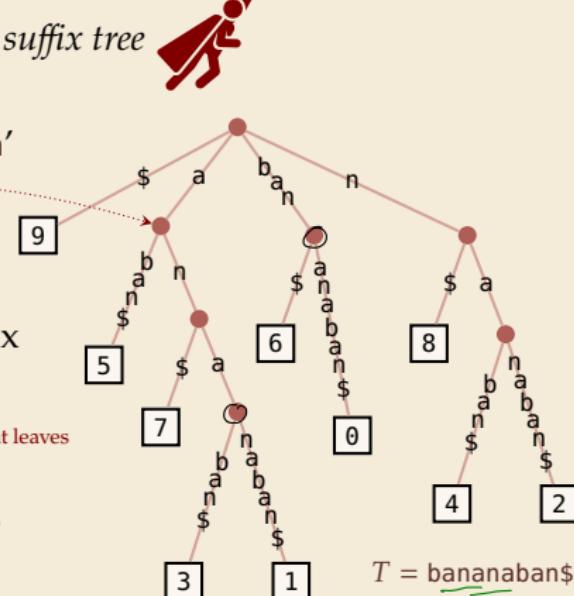
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- Algorithm:

1. Compute string depth (=length of path label) of nodes
2. Find internal nodes with maximal string depth

actually: adjacent leaves



- Both can be done in depth-first traversal  $\rightsquigarrow \Theta(n)$  time