

6 Text Indexing – Searching whole genomes

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Outline

6 Text Indexing

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6.1 Motivation

Text indexing

- ▶ *Text indexing* (also: *offline text search*):

- ▶ case of string matching: find $P[0..m-1]$ in $T[0..n-1]$
- ▶ but with *fixed* text \rightsquigarrow preprocess T (instead of P)
- \rightsquigarrow expect many queries P , answer them without looking at all of T
- \rightsquigarrow essentially a data structuring problem: “building an *index* of T ”

Latin: “one who points out”

- ▶ application areas

- ▶ web search engines
- ▶ online dictionaries
- ▶ online encyclopedia
- ▶ DNA/RNA data bases
- ▶ ... searching in any collection of text documents (that grows only moderately)

Inverted indices

same as "indexes"

- ▶ original indices in books: list of (key) words \mapsto page numbers where they occur
 - ▶ assumption: searches are only for **whole** (key) **words**
- \rightsquigarrow often reasonable for natural language text

Inverted index:

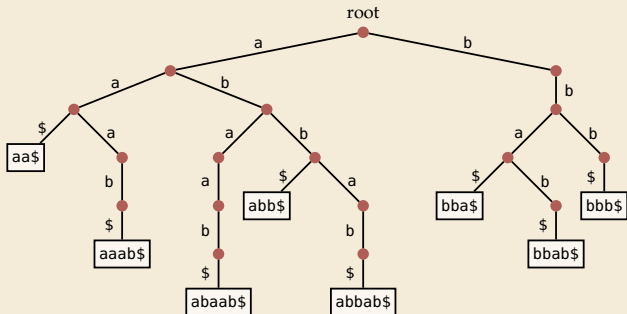
- ▶ collect all words in T
 - ▶ can be as simple as splitting T at whitespace
 - ▶ actual implementations typically support *stemming* of words
goes \rightarrow go, cats \rightarrow cat
- ▶ store mapping from words to a list of occurrences \rightsquigarrow *how?*

Tries

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

- ▶ **Example:**

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

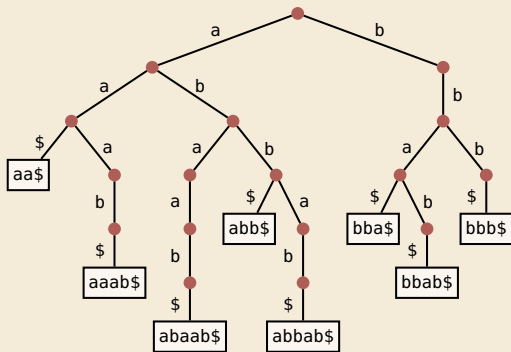


Compact tries

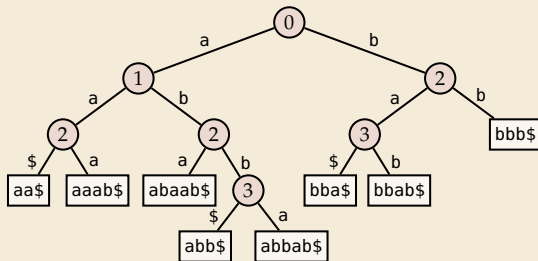
- ▶ compress paths of unary nodes into single edge
- ▶ nodes store index of next character

=1 child

standard trie



compact trie



~ searching slightly trickier, but same time complexity as in trie

- ▶ all nodes ≥ 2 children ~ $\#nodes \leq \#leaves = \#strings$ ~ linear space

Tries as inverted index

👍 simple

👍 fast lookup

👎 cannot handle more general queries:

- ▶ search part of a word
- ▶ search phrase (sequence of words)

👎 what if the 'text' does not even have words to begin with?!

- ▶ biological sequences

```
ACAAGATGCCATTGTCCCCCGGCTCCTGCTGCTGCTCTCCGGGGCCACGGCCACCGCTGCCCTGCCCTGGAGGGTGGCCCCACCGGC  
CGAGACAGCGAGCATATGCAGGAAGCGGCAGGAATAAGGAAAAGCAGCCTCCTGACTTTCCTCGCTTGGTGGTTTGAGTGGACCTCCAGGC  
CAGTGGCGGGCCCCCATAGGAGAGGAAGCTCGGGAGGTGGCCAGGCGGCAGGAAGGCGCACCCCCCAGCAATCCGCGCGCCGGGACAGAA  
TGCCCTGCAGGAACCTTCTTCTGGAAGACCTTCTCCTCTGCAAATAAACCTCACCCATGAATGCTCACGCAAGTTTAATTACAGACCTGAA
```

- ▶ binary streams

```
00000010101001111010111000001111100011111011111001101101000011100010011011110000010001101010  
0110110000110101101000000010000000011101011000001000011110101110110010001100101101110111111  
110001010001011001010000001110101010011000000001101100001100111110000101 0101011101111000011  
10101110010010101010100000111110100110000001111001101010000000100100100000101100011000110111
```

~> need new ideas

6.2 Suffix Trees

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

↪ alive



Can we do this in time $O(|S_1| + \dots + |S_k|)$? How??

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

- ▶ suffix tree \mathcal{T} for text $T = T[0..n - 1]$ = compact trie of all suffixes of $T\$$ (set $T[n] := \$$)
- ▶ except: in leaves, store *start index* (instead of actual string)

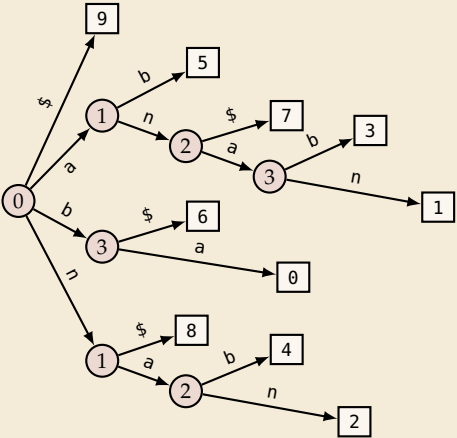
Example:

$T = \text{bananaban\$}$

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

0	1	2	3	4	5	6	7	8	9
b	a	n	a	n	a	b	a	n	\$

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



Suffix trees – Construction

- ▶ $T[0..n-1]$ 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. But that takes time $\Theta(n^2)$. \rightsquigarrow not interesting!



same order of growth as reading the text!

Amazing result: Can construct the suffix tree of T in $\Theta(n)$ time!

- ▶ algorithms are a bit tricky to understand
- ▶ but were a theoretical breakthrough
- ▶ and they are efficient in practice (and heavily used)!

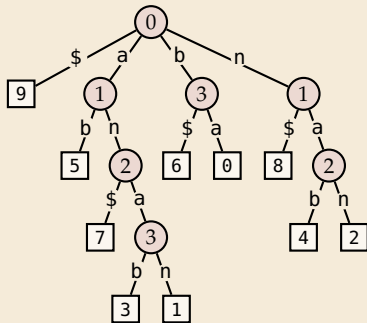
\rightsquigarrow for now, take linear-time construction for granted. What can we do with them?

6.3 Applications

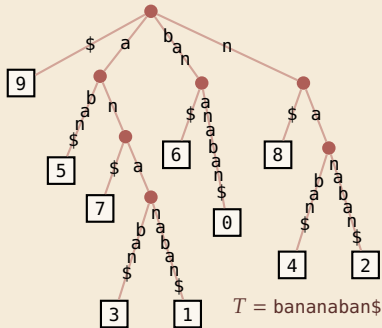
Applications of suffix trees

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

Recall: \mathcal{T} stored like this:



but think about this:



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

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*?



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

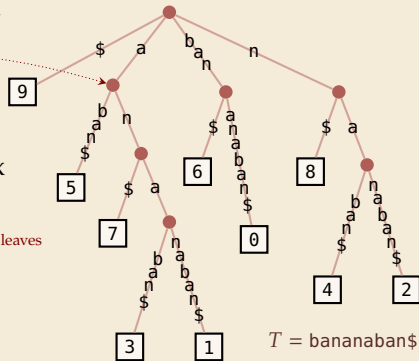
↪ longest repeated substring = longest common prefix (LCP) of two suffixes

actually: adjacent leaves

- Algorithm:

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

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



Generalized suffix trees

- ▶ longest *repeated* substring (of one string) feels very similar to longest *common* substring of several strings $T^{(1)}, \dots, T^{(k)}$ with $T^{(j)} \in \Sigma^{n_j}$
 - ▶ can we solve that in the same way?
 - ▶ could build the suffix tree for each $T^{(j)}$... but doesn't seem to help
- ↪ need a *single/joint* suffix tree for *several* texts

Enter: *generalized suffix tree*

- ▶ Define $T := T^{(1)}\$_1 T^{(2)}\$_2 \dots T^{(k)}\$_k$ for k new end-of-word symbols
- ▶ Construct suffix tree \mathcal{T} for T

↪ $\$_j$ -edges always leads to leaves ↪ \exists leaf (j, i) for each suffix $T_i^{(j)} = T^{(j)}[i..n_j]$



Application 3: Longest common substring

- ▶ With that new idea, we can find longest common superstrings:
 1. Compute generalized suffix tree \mathcal{T} .
 2. Store with each node the *subset of strings* that contain its path label:
 - 2.1. Traverse \mathcal{T} bottom-up.
 - 2.2. For a leaf (j, i) , the subset is $\{j\}$.
 - 2.3. For an internal node, the subset is the union of its children.
 3. In top-down traversal, compute *string depths* of nodes. (as above)
 4. Report deepest node (by string depth) whose subset is $\{1, \dots, k\}$.

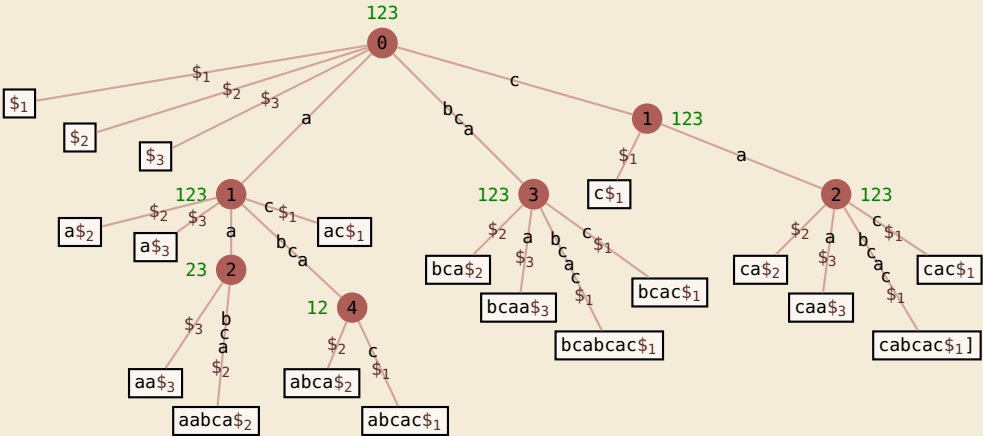
- ▶ Each step takes time $\Theta(n)$ for $n = n_1 + \dots + n_k$ the total length of all texts.

“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)]

Longest common substring – Example

$T^{(1)} = \text{bcabcac}$, $T^{(2)} = \text{aabca}$, $T^{(3)} = \text{bcaa}$



6.4 Longest Common Extensions

Application 4: Longest Common Extensions

- ▶ We implicitly used a special case of a more general, versatile idea:

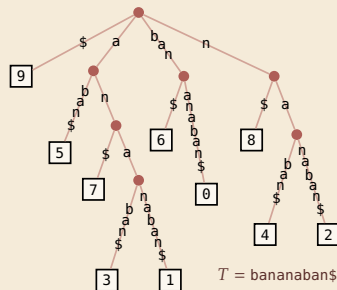
The *longest common extension (LCE)* data structure:

- ▶ **Given:** String $T[0..n-1]$
- ▶ **Goal:** Answer LCE queries, i.e.,
given positions i, j in T ,
how far can we read the same text from there?
formally: $\text{LCE}(i, j) = \max\{\ell : T[i..i+\ell] = T[j..j+\ell]\}$

↪ use suffix tree of T !

- ▶ In \mathcal{T} : $\text{LCE}(i, j) = \text{LCP}(T_i, T_j) \rightsquigarrow$ same thing, different name!
= string depth of
lowest common ancestor (LCA) of
leaves \boxed{i} and \boxed{j}

- ▶ in short: $\text{LCE}(i, j) = \text{LCP}(T_i, T_j) = \text{stringDepth}(\text{LCA}(\boxed{i}, \boxed{j}))$



Efficient LCA

How to find lowest common ancestors?

- ▶ Could walk up the tree to find LCA $\rightsquigarrow \Theta(n)$ worst case 🙄
- ▶ Could store all LCAs in big table $\rightsquigarrow \Theta(n^2)$ space and preprocessing 🙄



Amazing result: Can compute data structure in $\Theta(n)$ time and space that finds any LCA is **constant(!) time**.

- ▶ a bit tricky to understand
- ▶ but a theoretical breakthrough
- ▶ and useful in practice

and suffix tree construction inside ...



\rightsquigarrow for now, use $O(1)$ LCA as black box.

\rightsquigarrow After linear preprocessing (time & space), we can find LCEs in $O(1)$ time.

Application 5: Approximate matching

k-mismatch matching:

- ▶ **Input:** text $T[0..n-1]$, pattern $P[0..m-1]$, $k \in [0..m)$
- ▶ **Output:** “Hamming distance $\leq k$ ”
 - ▶ smallest i so that $T[i..i+m)$ are P differ in at most k characters
 - ▶ or NO_MATCH if there is no such i

↪ searching with typos

- ▶ Assume longest common extensions in $T\$_1P\$_2$ can be found in $O(1)$
 - ↪ generalized suffix tree \mathcal{T} has been built
 - ↪ string depths of all internal nodes have been computed
 - ↪ constant-time LCA data structure for \mathcal{T} has been built

Kangaroo Algorithm for approximate matching



```
1 procedure kMismatch( $T[0..n - 1]$ ,  $P[0..m - 1]$ )
2   // build LCE data structure
3   for  $i := 0, \dots, n - m - 1$  do
4     mismatches := 0;  $t := i$ ;  $p := 0$ 
5     while mismatches  $\leq k \wedge p < m$  do
6        $\ell := \text{LCE}(t, p)$  // jump over matching part
7        $t := t + \ell + 1$ ;  $p := p + \ell + 1$ 
8       mismatches := mismatches + 1
9     if  $p == m$  then
10      return  $i$ 
```

► **Analysis:** $\Theta(n + m)$ preprocessing + $O(n \cdot k)$ matching

\rightsquigarrow very efficient for small k

► State of the art

- $O(n \frac{k^2 \log k}{m})$ possible with complicated algorithms
- extensions for edit distance $\leq k$ possible

Application 6: Matching with wildcards

- ▶ Allow a wildcard character in pattern

stands for arbitrary (single) character

unit*	P
in_unit5_we_will	T

- ▶ similar algorithm as for k -mismatch $\rightsquigarrow O(n \cdot k + m)$ when P has k wildcards

* * *

Many more applications, in particular for problems on biological sequences

20+ described in Gusfield, *Algorithms on strings, trees, and sequences* (1999)

Suffix trees – Discussion

► Suffix trees were a threshold invention



linear time and space



suddenly many questions efficiently solvable in theory



construction of suffix trees:

linear time, but significant overhead



construction methods fairly complicated

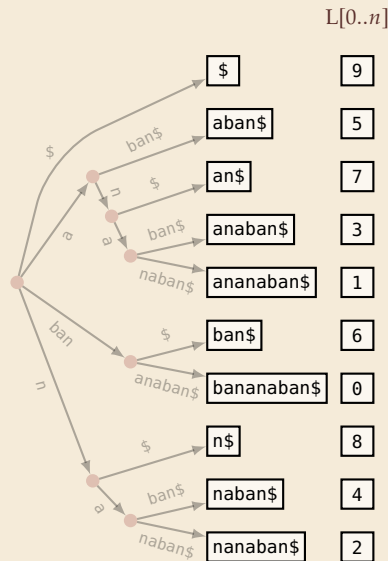


many pointers in tree incur large space overhead



6.5 Suffix Arrays

Putting suffix trees on a diet



► **Observation:** order of leaves in suffix tree
= suffixes lexicographically *sorted*

► Idea: only store list of leaves $L[0..n]$

► Enough to do efficient string matching!

1. Use binary search for pattern P

2. check if P is prefix of suffix after found position

► **Example:** $P = \text{ana}$

↪ $L[0..n]$ is called *suffix array*:

$L[r] = (\text{start index of } r\text{th suffix in sorted order})$

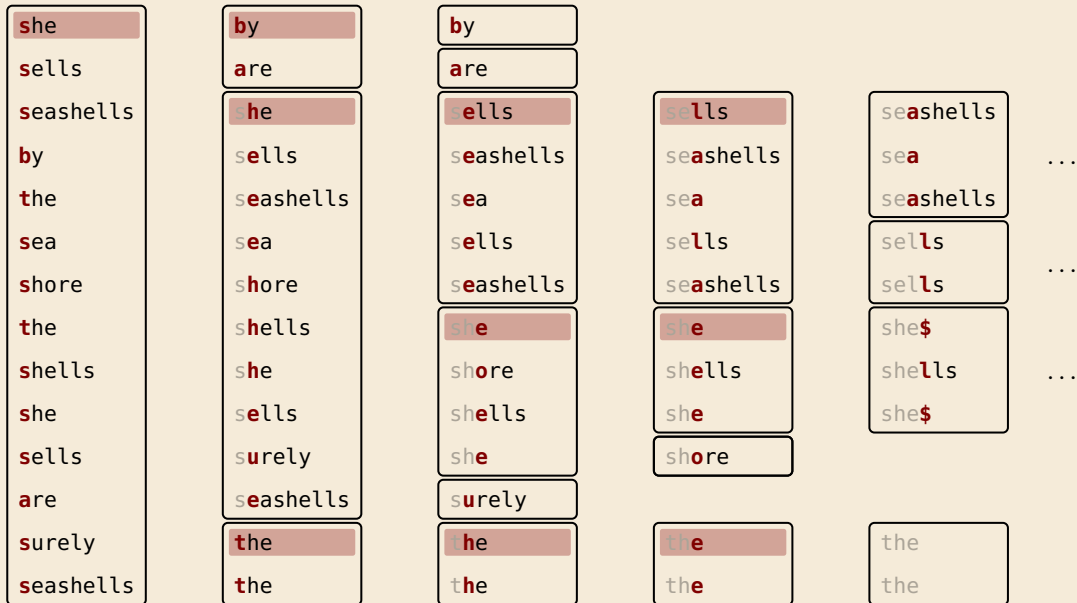
► using L , can do string matching with
 $\leq (\lg n + 2) \cdot m$ character comparisons

Suffix arrays – Construction

How to compute $L[0..n]$?

- ▶ from suffix tree
 - ▶ possible with traversal . . .
 - 👎 but we are trying to avoid constructing suffix trees!
- ▶ sorting the suffixes of T using general purpose sort
 - 👍 trivial to code!
 - ▶ but: comparing two suffixes can take $\Theta(n)$ character comparisons
 - 👎 $\Theta(n^2 \log n)$ time in worst case
- ▶ we do better!

Fat-pivot radix quicksort – Example



Fat-pivot radix quicksort

details in §5.1 of Sedgewick, Wayne *Algorithms 4th ed.* (2011), Pearson

► **partition** based on *d*th character only (initially $d = 0$)

↪ 3 segments: smaller, equal, or larger than *d*th symbol of pivot

► recurse on smaller and large with same *d*, on equal with $d + 1$

↪ never compare equal prefixes twice

↪ can show: $\sim 2 \ln(2) \cdot n \lg n \approx 1.39n \lg n$ character comparisons in expectation

👍 simple to code

👍 efficient for sorting many lists of strings

► fat-pivot radix quicksort finds suffix array in $O(n \log n)$ expected time

random pivots



but we can do $O(n)$ time worst case!

6.6 Linear-Time Suffix Sorting

Inverse suffix array: going left & right

► to understand the fastest algorithm, it is helpful to define the *inverse suffix array*:

- $R[i] = r \iff L[r] = i$ $L = \text{leaf array}$
 - \iff there are r suffixes that come before T_i in sorted order
 - $\iff T_i$ has (0-based) *rank* $r \rightsquigarrow$ call $R[0..n]$ the *rank array*

i	$R[i]$	T_i		r	$L[r]$	$T_{L[r]}$
0	6 th	bananaban\$		0	9	\$
1	4 th	ananaban\$		1	5	aban\$
2	9 th	nanaban\$		2	7	an\$
3	3 th	anaban\$		3	3	anaban\$
4	8 th	naban\$		4	1	ananaban\$
5	1 th	aban\$		5	6	ban\$
6	5 th	ban\$		6	0	bananaban\$
7	2 th	an\$		7	8	n\$
8	7 th	n\$		8	4	naban\$
9	0 th	\$		9	2	nanaban\$

right
 $R[0] = 6$

left
 $L[8] = 4$

sort suffixes

Linear-time suffix sorting

DC3 / Skew algorithm

1. Compute rank array $R_{1,2}$ for suffixes T_i starting at $i \not\equiv 0 \pmod{3}$ *not a multiple of 3* \downarrow *recursively*.
2. Induce rank array R_3 for suffixes $T_0, T_3, T_6, T_9, \dots$ from $R_{1,2}$.
3. Merge $R_{1,2}$ and R_3 using $R_{1,2}$.
 \rightsquigarrow rank array R for entire input

► We will show that steps 2. and 3. take $\Theta(n)$ time

\rightsquigarrow Total complexity is $n + \frac{2}{3}n + \left(\frac{2}{3}\right)^2 n + \left(\frac{2}{3}\right)^3 n + \dots \leq n \cdot \sum_{i \geq 0} \left(\frac{2}{3}\right)^i = 3n = \Theta(n)$

► **Note:** L can easily be computed from R in one pass, and vice versa.

\rightsquigarrow Can use whichever is more convenient.

DC3 / Skew algorithm – Step 2: Inducing ranks

- ▶ **Assume:** rank array $R_{1,2}$ known:

$$\text{▶ } R_{1,2}[i] = \begin{cases} \text{rank of } T_i \text{ among } T_1, T_2, T_4, T_5, T_7, T_8, \dots & \text{for } i = 1, 2, 4, 5, 7, 8, \dots \\ \text{undefined} & \text{for } i = 0, 3, 6, 9, \dots \end{cases}$$

- ▶ **Task:** sort the suffixes $T_0, T_3, T_6, T_9, \dots$ in linear time (!)

- ▶ Suppose we want to compare T_0 and T_3 .

- ▶ Characterwise comparisons too expensive
- ▶ but: after removing first character, we obtain T_1 and T_4
- ▶ these two can be compared in *constant time* by comparing $R_{1,2}[1]$ and $R_{1,2}[4]$!

~> T_0 comes before T_3 in lexicographic order
iff pair $(T[0], R_{1,2}[1])$ comes before pair $(T[3], R_{1,2}[4])$ in lexicographic order

DC3 / Skew algorithm – Inducing ranks example

$T = \text{hannahbansbananasman}\$ \$ \$$

(append 3 \$ markers)

T_0 hannahbansbananasman\$\$\$
 T_3 nahbansbananasman\$\$\$
 T_6 bansbananasman\$\$\$
 T_9 sbananasman\$\$\$
 T_{12} nanasman\$\$\$
 T_{15} asman\$\$\$
 T_{18} an\$\$\$
 T_{21} \$\$

$\text{sman}\$ \$ \$ = T_{16}$

T_0 h05
 T_3 n02
 T_6 b06
 T_9 s07
 T_{12} n04
 T_{15} a14
 T_{18} a10
 T_{21} \$00

$R_{1,2}[16] = 14$

T_1 annahbansbananasman\$\$\$
 T_2 nnahbansbananasman\$\$\$
 T_4 ahbansbananasman\$\$\$
 T_5 hbansbananasman\$\$\$
 T_7 ansbananasman\$\$\$
 T_8 nsbananasman\$\$\$
 T_{10} bananasman\$\$\$
 T_{11} ananasman\$\$\$
 T_{13} anasman\$\$\$
 T_{14} nasman\$\$\$
 T_{16} sman\$\$\$
 T_{17} man\$\$\$
 T_{19} n\$\$\$
 T_{20} \$\$\$
 T_{22} \$

$R_{1,2}[22] = 0$
 $R_{1,2}[20] = 1$
 $R_{1,2}[4] = 2$
 $R_{1,2}[11] = 3$
 $R_{1,2}[13] = 4$
 $R_{1,2}[1] = 5$
 $R_{1,2}[7] = 6$
 $R_{1,2}[10] = 7$
 $R_{1,2}[5] = 8$
 $R_{1,2}[17] = 9$
 $R_{1,2}[19] = 10$
 $R_{1,2}[14] = 11$
 $R_{1,2}[2] = 12$
 $R_{1,2}[8] = 13$
 $R_{1,2}[16] = 14$

T_{22} \$
 T_{20} \$\$\$
 T_4 ahbansbananasman\$\$\$
 T_{11} ananasman\$\$\$
 T_{13} anasman\$\$\$
 T_1 annahbansbananasman\$\$\$
 T_7 ansbananasman\$\$\$
 T_{10} bananasman\$\$\$
 T_5 hbansbananasman\$\$\$
 T_{17} man\$\$\$
 T_{19} n\$\$\$
 T_{14} nasman\$\$\$
 T_2 nnahbansbananasman\$\$\$
 T_8 nsbananasman\$\$\$
 T_{16} sman\$\$\$

$R_{1,2}$ (known)

radix sort

T_{21} \$00 $\rightsquigarrow R_0[21] = 0$
 T_{18} a10 $\rightsquigarrow R_0[18] = 1$
 T_{15} a14 $\rightsquigarrow R_0[15] = 2$
 T_6 b06 $\rightsquigarrow R_0[6] = 3$
 T_0 h05 $\rightsquigarrow R_0[0] = 4$
 T_3 n02 $\rightsquigarrow R_0[3] = 5$
 T_{12} n04 $\rightsquigarrow R_0[12] = 6$
 T_9 s07 $\rightsquigarrow R_0[9] = 7$
 R_0

► sorting of pairs doable in $O(n)$ time by 2 iterations of counting sort

\rightsquigarrow Obtain R_0 in $O(n)$ time

DC3 / Skew algorithm – Step 3: Merging

T_{21} \$\$
 T_{18} an\$\$\$
 T_{15} asman\$\$\$
 T_6 bansbananasman\$\$\$
 T_0 hannahbansbananasman\$\$\$
 T_3 nahbansbananasman\$\$\$
 T_{12} nanasman\$\$\$
 T_9 sbananasman\$\$\$

T_{22} \$
 T_{20} \$\$\$
 T_4 ahbansbananasman\$\$\$
 T_{11} ananasman\$\$\$
 T_{13} anasman\$\$\$
 T_1 annahbansbananasman\$\$\$
 T_7 ansbananasman\$\$\$
 T_{10} bananasman\$\$\$
 T_5 hbansbananasman\$\$\$
 T_{17} man\$\$\$
 T_{19} n\$\$\$
 T_{14} nasman\$\$\$
 T_2 nnahbansbananasman\$\$\$
 T_8 nsbananasman\$\$\$
 T_{16} sman\$\$\$

T_{22} \$
 T_{21} \$\$
 T_{20} \$\$\$
 T_4 ahbansbananasman\$\$\$
 T_{18} an\$\$\$

► Have:

► sorted 1,2-list:

$T_1, T_2, T_4, T_5, T_7, T_8, T_{10}, T_{11}, \dots$

► sorted 0-list:

$T_0, T_3, T_6, T_9, \dots$

► Task: Merge them!

► use standard merging method from Mergesort

► but speed up comparisons using $R_{1,2}$

$\leadsto O(n)$ time for merge

Compare T_{15} to T_{11}

Idea: try same trick as before

$T_{15} = \text{asman}$$$
 $= \text{asman}$$$ can't compare T_{16}
 $= aT_{16}$ and T_{12} either!
 $T_{11} = \text{ananasman}$$$
 $= \text{ananasman}$$$
 $= aT_{12}$$$$$

\leadsto Compare T_{16} to T_{12}

$T_{16} = \text{sman}$$$
 $= \text{sman}$$$ always at most 2 steps
 $= sT_{17}$ then can use $R_{1,2}$!
 $T_{12} = \text{nanasman}$$$
 $= \text{aanasman}$$$
 $= aT_{13}$$$$$

DC3 / Skew algorithm – Fix recursive call

- ▶ both step 2. and 3. doable in $O(n)$ time!
 - ▶ But: we cheated in 1. step! “compute rank array $R_{1,2}$ recursively”
 - ▶ Taking a *subset* of suffixes is *not* an instance of the same problem!
- ↪ Need a single *string* T' to recurse on, from which we can deduce $R_{1,2}$.



How can we make T' “skip” some suffixes?



redefine alphabet to be *triples of characters* \boxed{abc}

↪ suffixes of $T^\square \iff T_0, T_3, T_6, T_9, \dots$

▶ $T' = T[1..n)^\square \boxed{\$ \$ \$} T[2..n)^\square \boxed{\$ \$ \$} \iff T_i$ with $i \not\equiv 0 \pmod{3}$.

↪ Can call suffix sorting recursively on T' and map result to $R_{1,2}$

$T = \text{bananaban} \$ \$ \$$
↪ $T^\square = \boxed{\text{ban}} \boxed{\text{ana}} \boxed{\text{ban}} \boxed{\$ \$ \$}$
 $\boxed{\text{ana}} \boxed{\text{ban}} \boxed{\$ \$ \$}$
 $\boxed{\text{ban}} \boxed{\$ \$ \$}$
 $\boxed{\$ \$ \$}$

DC3 / Skew algorithm – Fix alphabet explosion

► Still does not quite work!

► Each recursive step *cubes* σ by using triples!

↪ (Eventually) cannot use linear-time sorting anymore!

► But: Have at most $\frac{2}{3}n$ different triples \boxed{abc} in T' !

↪ Before recursion:

1. Sort all occurring triples. (using counting sort in $O(n)$)

2. Replace them by their *rank* (in Σ).

↪ Maintains $\sigma \leq n$ without affecting order of suffixes.

DC3 / Skew algorithm – Step 3. Example

$$T' = T[1..n) \square \square \square T[2..n) \square \square \square$$

► $T = \text{hannahbansbananasman\$}$ $T_2 = \text{nnaahbansbananasman\$}$

$T' =$ annahbansbananasman\$\$ \$\$\$ nnahbansbananasman\$\$\$

► Occurring triples:

annahbansbananasman\$\$ \$\$\$ nnahbansb nasman

► Sorted triples with ranks:

Rank	00	01	02	03	04	05	06	07	08	09	10	11	12
Triple	\$\$\$	ahb	ana	ann	ans	ban	hba	man	n\$\$	nas	nna	nsb	sma

► $T' =$ annahbansbananasman\$\$ \$\$\$ nnahbansbananasman\$\$\$

$T'' =$ 03 01 04 05 02 12 08 00 10 06 11 02 09 07 00

Suffix array – Discussion

- 👍 sleek data structure compared to suffix tree
- 👍 simple and fast $O(n \log n)$ construction
- 👍 more involved but fast $O(n)$ construction
- 👍 supports efficient string matching
- 👎 string matching takes $O(m \log n)$, not optimal $O(m)$
- 👎 Cannot use more advanced suffix tree features
e. g., for longest repeated substrings



6.7 The LCP Array

String depths of internal nodes

- Recall algorithm for longest repeated substring in **suffix tree**

1. Compute *string depth* of nodes
2. Find *path label* to node with maximal string depth

- Can we do this using **suffix arrays**?

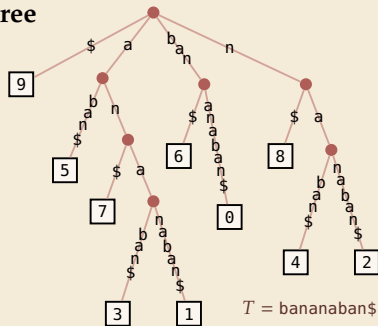
- Yes, by **enhancing** the suffix array with the **LCP array**!

$\text{LCP}[1..n]$

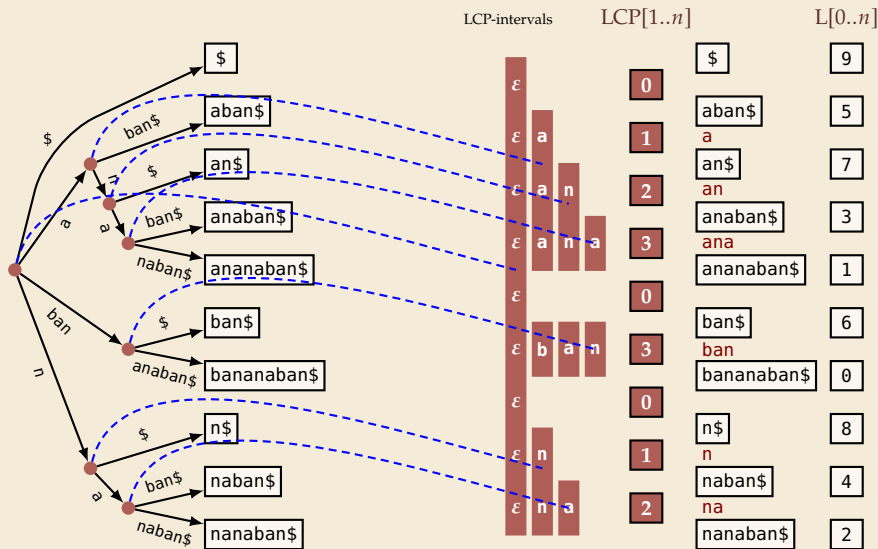
$\text{LCP}[r] = \text{LCP}(T_{L[r]}, T_{L[r-1]})$

longest common prefix of suffixes of rank r and $r - 1$

\rightsquigarrow longest repeated substring = find maximum in $\text{LCP}[1..n]$



LCP array and internal nodes




↪ Leaf array $L[0..n]$ plus LCP array $LCP[1..n]$ encode full tree!

LCP array construction

- ▶ computing $LCP[1..n]$ naively too expensive

- ▶ each value could take $\Theta(n)$ time

-  $\Theta(n^2)$ in total

- ▶ but: seeing one large (= costly) LCP value \rightsquigarrow can find another large one!

- ▶ Example: $T = \text{Buffalo_buffalo_buffalo_buffalo\$}$

- ▶ first few suffixes in sorted order:

$T_{L[0]} = \$$

$T_{L[1]} = \text{alo_buffalo\$}$

$T_{L[2]} = \text{alo_buffalo_buffalo\$}$

alo_buffalo_buffalo $\rightsquigarrow LCP[3] = 19$

$T_{L[3]} = \text{alo_buffalo_buffalo_buffalo\$}$

- \rightsquigarrow **Removing first character** from $T_{L[2]}$ and $T_{L[3]}$ gives two new suffixes:

$T_{L[?]} = \text{lo_buffalo_buffalo\$}$

lo_buffalo_buffalo $\rightsquigarrow LCP[?] = 18$

$T_{L[?]} = \text{lo_buffalo_buffalo_buffalo\$}$

\uparrow
unclear where...



Shortened suffixes might *not* be *adjacent* in sorted order!

\rightsquigarrow no LCP entry for them!

Kasai's algorithm – Example

- ▶ Kasai et al. used above observation systematically
- ▶ Key idea: *compute* LCP values in *text order*
- ▶ Dropping first character of adjacent suffixes might not lead to *adjacent* shorter suffixes, but LCP entry can only be *longer*.

i	$R[i]$	T_i	r	$L[r]$	$T_{L[r]}$	$LCP[r]$
0	6 th	bananaban\$	0	9	\$	–
1	4 th	ananaban\$	1	5	aban\$	0
2	9 th	nanaban\$	2	7	an\$	1
3	3 th	anaban\$	3	3	anaban\$	2
4	8 th	naban\$	4	1	ananaban\$	3
5	1 th	aban\$	5	6	ban\$	0
6	5 th	ban\$	6	0	bananaban\$	3
7	2 th	an\$	7	8	n\$	0
8	7 th	n\$	8	4	naban\$	1
9	0 th	\$	9	2	nanaban\$	2

Kasai's algorithm – Code

```
1 procedure computeLCP( $T[0..n]$ ,  $L[0..n]$ ,  $R[0..n]$ )
2   // Assume  $T[n] = \$$ ,  $L$  and  $R$  are suffix array and inverse
3    $\ell := 0$ 
4   for  $i := 0, \dots, n - 1$ 
5      $r := R[i]$ 
6     // compute LCP[ $r$ ]; note that  $r > 0$  since  $R[n] = 0$ 
7      $i_{-1} := L[r - 1]$ 
8     while  $T[i + \ell] == T[i_{-1} + \ell]$  do
9        $\ell := \ell + 1$ 
10    LCP[ $r$ ] :=  $\ell$ 
11     $\ell := \max\{\ell - 1, 0\}$ 
12 return LCP[ $1..n$ ]
```

- ▶ remember length ℓ of induced common prefix
- ▶ use L to get start index of suffixes

Analysis:

- ▶ dominant operation:
character comparisons
- ▶ separately count those with
outcomes “=” resp. “≠”
- ▶ each ≠ ends iteration of for-loop
 $\rightsquigarrow \leq n$ cmps
- ▶ each = implies increment of ℓ ,
but $\ell \leq n$ and
decremented $\leq n$ times
 $\rightsquigarrow \leq 2n$ cmps

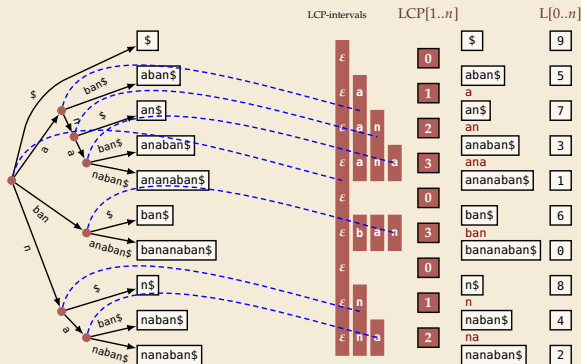
$\rightsquigarrow \Theta(n)$ overall time

Back to suffix trees

We can finally look into the black box of linear-time suffix-array construction!



1. Compute suffix array for T .
2. Compute LCP array for T .
3. Construct \mathcal{T} from suffix array and LCP array.



Conclusion

► *(Enhanced) Suffix Arrays* are the modern version of suffix trees

👎 can be harder to reason about

👍 can support same algorithms as suffix trees

👍 but use much less space

👍 simpler linear-time construction