

### 3. Exact Paterson Matching

# String Matching

Given: pattern  $P$ , text  $T$  ( $P, T$  are strings)

Aim: find occurrences of  $P$  in  $T$

Exmpl:  $T =$  <sup>1 2 3 4 5 6 7 8 9 10 ...</sup> AATGCATGCA .....  
 $P =$  ATG

$P$  occurs on position 2, 6, ...

This has applications in:

- Bioinformatics (eg. sequence assembly where one may align fragments of your DNA to reference genome to get read of your DNA; also applications in finding repeated regions & many more)
- general word processing
- internet search
- "fgrep" in Unix
- search for plagiarism
- subtask for e.g. inexact matching

## Basics:

string  $S = x_1 \dots x_n$ ,  $|S| = n$

$$S[i..j] = x_i x_{i+1} \dots x_j$$

$$S(i) = x_i$$

$$S[1..j] \hat{=} \text{prefix of } S \text{ ending at } j$$

$$S[j..n] \hat{=} \text{suffix of } S \text{ starting at } j$$

S: <sup>1 2 3 4 5 6 7 8</sup> HONOLULU  
HONO (1..4) prefix  
LU (7..8) suffix  
 $S(S) = L$

if  $P(i) = T(k)$  for some  $i, k$  then they match  
else mismatch.

## Naive Method

NAIVE (P, T)

// compare  $T(i \dots i + |P| - 1)$   
// with P for all  $i = 1 \dots |T| - |P| + 1$

Occurrences =  $\emptyset$

FOR (  $i = 1 \dots |T| - |P| + 1$  )

// loop over "pos." in T

FOR (  $j = 1 \dots |P|$  )

// loop over P

match = TRUE

IF (  $P(j) \neq T(i+j-1)$  )

// compare characters

match = FALSE

break

IF (match)

add  $i$  to occurrences

// save pos. on which

P occurs

RETURN occurrences.

Q: How often can P occur in T?

A:  $|T| - |P| + 1$  times

T = A A A A A A

P = A A

P occurs at pos. 1, 2, 3, 4, 5

= 5 times =  $6 - 2 + 1$

Q: What is greatest nr of character comparisons?

A:  $|P| \cdot (|T| - |P| + 1)$  (worst case)

See example above:  $2 \cdot 5 = 10$  comparisons

Q: What is least nr of character comparisons?

A:  $|T| - |P| + 1$  (best case)

T = A A A A A A

P = B A

⇒ RUNTIME NAIVE-alg

$|T| - |P| + 1$

$|P|$

1. loop

2 loop

$$\Rightarrow |T|(|P| - |P|^2 + 1) \leq |T||P|$$

$$|T| \geq |P|$$

⇒  $O(|T||P|)$  time

look now to one of many linear-time algorithms, i.e. instead of  $O(|P||T|)$   
we have runtime  $O(|P| + |T|)$

Z - algorithm [fundamental preprocessing used in many alg]

General idea: pre-process  $P$  in  $O(|P|)$  time to gain insight of internal structure of  $P$

DEF: let  $S$  be a string (usually  $S = P$  position)  
&  $i > 1$ .

$z_i := z_i(S) =$  length of longest substring in  $S$  that starts at position  $i$  & matches prefix of  $S$

Exmpl:

$i$	1	2	3	4	5	6	7	8	9	10	11
$S$	a	a	b	c	a	a	b	x	a	a	z
$z_i$	-	1	0	0	3	1	0	0	2	1	0

DEF (z-Box):  $\forall i > 1$  s.t.  $z_i > 0$ , the z-Box at  $i$  is the interval  $[i, i + z_i - 1]$

Exmpl:

$i$	1	2	3	4	5	6	7	8	9	10	11
$S$	a	a	b	c	a	a	b	x	a	a	z
$z_i$	-	1	0	0	3	1	0	0	2	1	0

$z_i > 0$  at pos. 2, 5, 6, 9, 10

$$\begin{aligned} z\text{-Box at } 2 &= [2, 2+1-1] = [2, 2] \\ \text{at } 5 &= [5, 5+3-1] = [5, 7] \\ \text{at } 6 &= [6, 6+1-1] = [6, 6] \\ \text{at } 9 &= [9, 9+2-1] = [9, 10] \\ \text{at } 10 &= [10, 10+1-1] = [10, 10] \end{aligned}$$

$i$	1	2	3	4	5	6	7	8	9	10	11
$S$	a	a	b	c	a	a	b	x	a	a	z

*(Red brackets in the original image group the 'a's at positions 2, 5-6, and 9-10.)*

These intervals  $[i, j]$  correspond  
to  $S[i..j]$  = longest substring  
starting at  $i \geq 1$  &  
matches prefix of  $S$

DEF

$\forall i > 1 :$

(r = right)

$r_i$  denotes right-most endpoint of z-Boxes  $z_j$  with  $z_j > 0$  &  $j \leq i$

(equ. to  $r_i =$  largest value of  $j + z_j - 1$  over all  $1 \leq j \leq i$  st  $z_j > 0$ )

"store index  $j$ ":  $l_i = j$  for  $j$  satisfying this  $\uparrow$

(l = left)

(equ. to  $l_i =$  position of left-end of z-Box ending in  $r_i$ )

if MORE than one z-Box ends in  $r_i$  then  $l_i$  can be chosen arbitrarily among those values

Formal:  $r_i = \max_{2 \leq j \leq i} \{ j + z_j - 1 : z_j > 0 \}$

$i$	1	2	3	4	5	6	7	8	9	10	11
S	a	a	b	c	a	a	b	x	a	a	z
$r_i$	-	2	2	2	7	7	7	7	10	10	10
$l_i$	-	2	2	2	5	5	5	5	9	9	9

right:  $r_6 = \max \left\{ \underbrace{2 + 1 - 1}_{\text{for } j=2}, \underbrace{5 + 3 - 1}_{j=5}, \underbrace{6 + 1 - 1}_{j=6} \right\} = 7$

left:  $l_6 = 5$

this essentially gives pos. of "large" z-Boxes.

## How to compute $z_i, r_i, l_i$ efficiently?

main idea: iterative approach

start with  $z_2$  (explicit comparison from left-to-right)

Assume  $\forall i < k$  values  $z_i, r_i, l_i$  have been computed.

for  $k$  use:  $z_i, l_i, r_i$  ( $i < k$ )

WITHOUT EXPLICIT character comparisons  
as much as possible.

## Overview of steps of $z$ -alg

① compute  $z_2$  (explicit comparison of  $S[1..n]$  with  $S[2..n]$  until mismatch)

IF ( $z_2 > 0$ )

ELSE

put  $l_2 = 2, r_2 = z_2 + 1$

put  $l_2 = r_2 = 0$

②

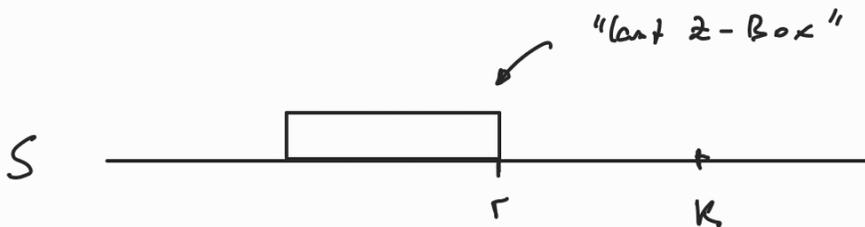
$k$ -th step ( $k \geq 2$ ).

$\Rightarrow$  all  $z_i, l_i, r_i$  computed for all  $i \leq k-1$

Compute  $z_k, l_k, r_k$  based on the following cases.

$$l := l_{k-1}, \quad r := r_{k-1}$$

Case 1:  $r < k$



in this case compute  $z_k$  via explicit comparison of  $S[k..n]$  &  $S[1..r]$  until mismatch.

IF ( $z_k > 0$ ) put  $l_k = k, r_k = k + z_k - 1$   
ELSE  $l_k = l, r_k = r$

Exmpl:

$i$	1	2	3	4	5	6	7	8	9	10	11
$S$	a	a	b	c	a	a	b	x	a	a	z
$r_i$	-	2	2	2	7	7	7	7			
$l_i$	-	2	2	2	5	5	5	5			

$k=9 \rightarrow r_g = 8 < k=9$  need to compare  $S[9..11]$  with  $S[1..n]$

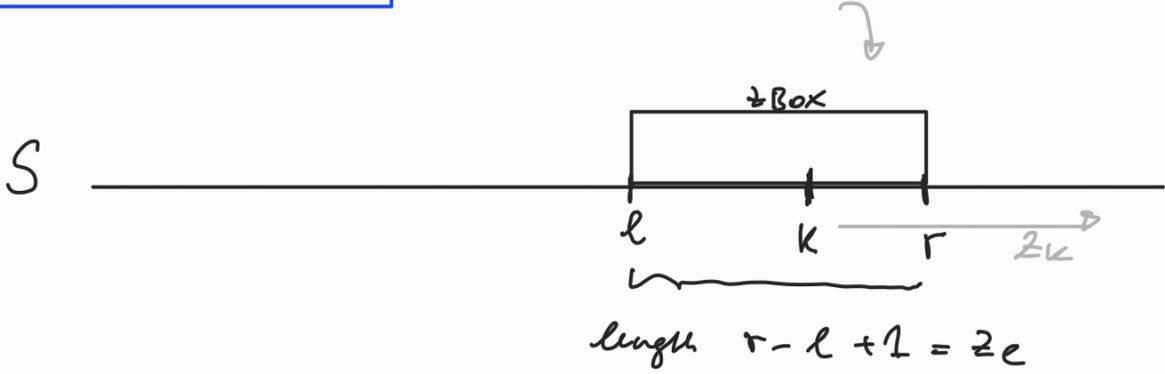
$\Rightarrow S[9] = S[1] \checkmark$   
 $S[10] = S[2] \checkmark$   
 $S[11] \neq S[3] \times$

$\Rightarrow z_g = 2, l_g = 9$

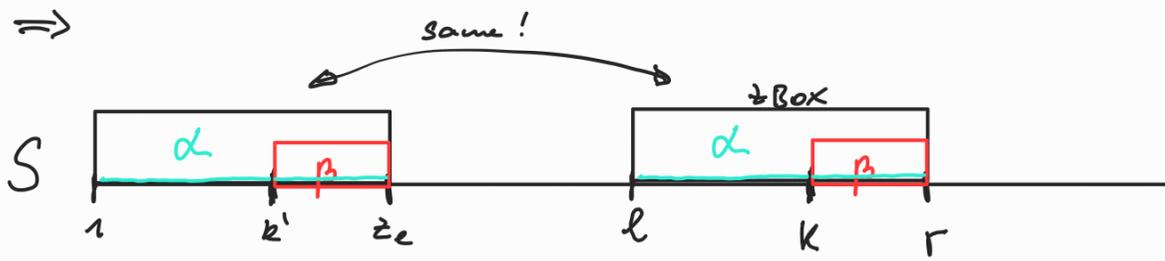
$r_g = 9 + 1 - 1 = 10$

Case 2:  $r \geq k$

[Now we have some pre-knowledge about about  $z_k$ ]



$z\text{-Box} \hat{=} \text{prefix of } S$



$$S[1..z_e] = S[l..r] = \alpha$$

$$S[k'..z_e] = S[k..r] = \beta$$

We want to know now  $z_k$  (longest string starting at  $k$  & is prefix of  $S$ )

Question: what we know about  $\beta$ ?

if we know  $\beta$  or "first part" of  $\beta$  is prefix of  $S$ , we don't need to compare the respective positions.

! This knowledge is already stored in  $z_{k'}$

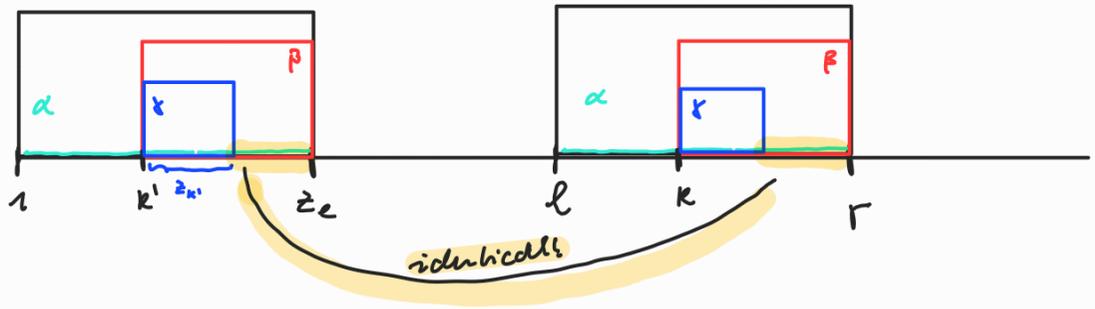
$$k' = k - l + 1$$

and leads to following subcases.

2A, 2B

2A

$$z_{k'} < |\beta| = r - k + 1$$



$z_{k'}$  = length of  $\gamma$  = length of longest substring starting at  $k'$  & is prefix of  $S$

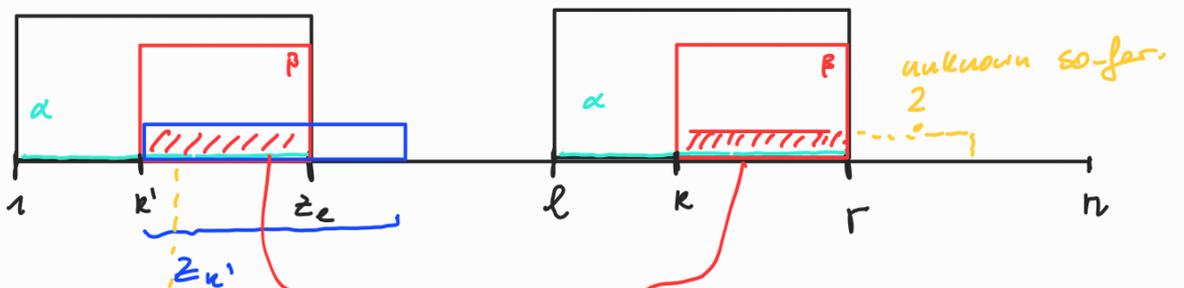
Since  $\beta = \beta$   
 left right  $\Rightarrow \gamma$  (right) is longest substring starting at  $k$  & is prefix of  $S$ .

$$\Rightarrow z_k = z_{k'}$$

$l$  &  $r$  remain unchanged.

2B

$$z_{k'} \geq |\beta| = r - k + 1$$



$\beta + 1$   
 pos.

this part matches already,  
 i.e.  $\beta$  (right) is part of longest  
 substring starting at  $k$  & prefix  
 of  $S$ .

$\beta = S[k..r]$  is prefix of  $S$ .

$$\text{so } z_k \geq |\beta| = r - k + 1$$

However  $Z_k > |S|$  may be possible.

$\Rightarrow$  compare  $S[r+1..n]$  <sup>[yellow part above]</sup>  
 with  $S[\beta|+1, \dots, n]$  until mismatch  
 //  $r - k + 1 + 1$   
 $- r - k + 2$   
 As possibly extended  
 Z-Box starting at  $k$

let  $q$  be pos. in  $S$  of 1st mismatch  
 & if no mismatch put  $q = |S| + 1$

put  $Z_k = q - k$  ( $= k + Z_k - 1$ )  
 $r = q - 1$   
 $l = k$

## Z Algorithm

```

1:  $r \leftarrow l \leftarrow 0$ 
2: for  $k = 2$  to  $|S|$  do
3: // Case 1:
4:   if  $r < k$  then
5:     Compare characters in  $S[k..n]$  with the ones in  $S[1..n]$  until mismatch is found (from left-to-right).
6:     Set  $Z_k$  to the length of the matched characters
7:     if  $Z_k > 0$  then
8:       Set  $r \leftarrow k + Z_k - 1, l \leftarrow k$ .
9: // Case 2:
10:  else
11:     $k' \leftarrow k - l + 1$ 
12: // Case 2a:
13:   if  $Z_{k'} < r - k + 1$  then
14:      $Z_k \leftarrow Z_{k'}$ 
15: // Case 2b:
16:   else
17:      $l \leftarrow k$ 
18:   Compare characters in  $S[r + 1..n]$  with the ones in  $S[r - k + 2..n]$  until mismatch is found
    (from left-to-right).
19:   let  $q > r$  be the position of the first mismatch or  $q = |S| + 1$  if no mismatch occurs
20:    $Z_k \leftarrow q - k, r \leftarrow q - 1$ 
  
```

later discussion implies:

Thm:  $Z$ -alg. correctly computes all  
 $Z + \text{Box } Z_i, i > 2$

however more important:

Thm: All  $Z_i(S)$  values are computed in  $O(|S|)$  time.

proof:

For loop (line 2-20) runs  $O(|S|)$  times.

Q: What happens within FOR-loop?

to answer this question let us count  
character-comparisons.

Each character comparison results either in match  
or mismatch.

$\Rightarrow$  let us count match / mismatches.

each comparison ends when 1st mismatch occurs

$\Rightarrow$  since  $O(|S|)$  different comparisons

$\Rightarrow$  total  $O(|S|)$  mismatches

Notation:  $C_k =$  NR of comparisons in  $k$ -th iteration  
 $m_k =$  NR of matches in  $k$ -th iteration

Claim:  $r_k - r_{k-1} \geq m_k$

Proof: in 2-Alg we either use Case 1/2A/2B.

Assume CASE 1 applies:

& thus,  $r_{k-1} < k$   $\xrightarrow{\text{Alg.}}$  explicit comparison of  $S[k..n]$  with  $S[1..m]$  until mismatch.

$\Rightarrow$  at most  $n-k+1$  comparisons

in Alg we put  $z_k = m_k$

in Alg case  $z_k > 0$  & [ $z_k = 0$  implicit by leaving " $r = r_k = r_{k-1}$ " unchanged].

$$z_k > 0 : r_k = k + z_k - 1 = k + m_k - 1 \geq r_{k-1} + m_k$$

$$\Leftrightarrow r_k \geq r_{k-1} + m_k$$

$$\Leftrightarrow r_k - r_{k-1} \geq m_k \quad \checkmark$$

$$z_k = 0 : r_k = r_{k-1} = r_{k-1} + m_k$$

$$\Leftrightarrow r_k - r_{k-1} = m_k = 0. \quad \checkmark$$

Now Case 2A/2B.

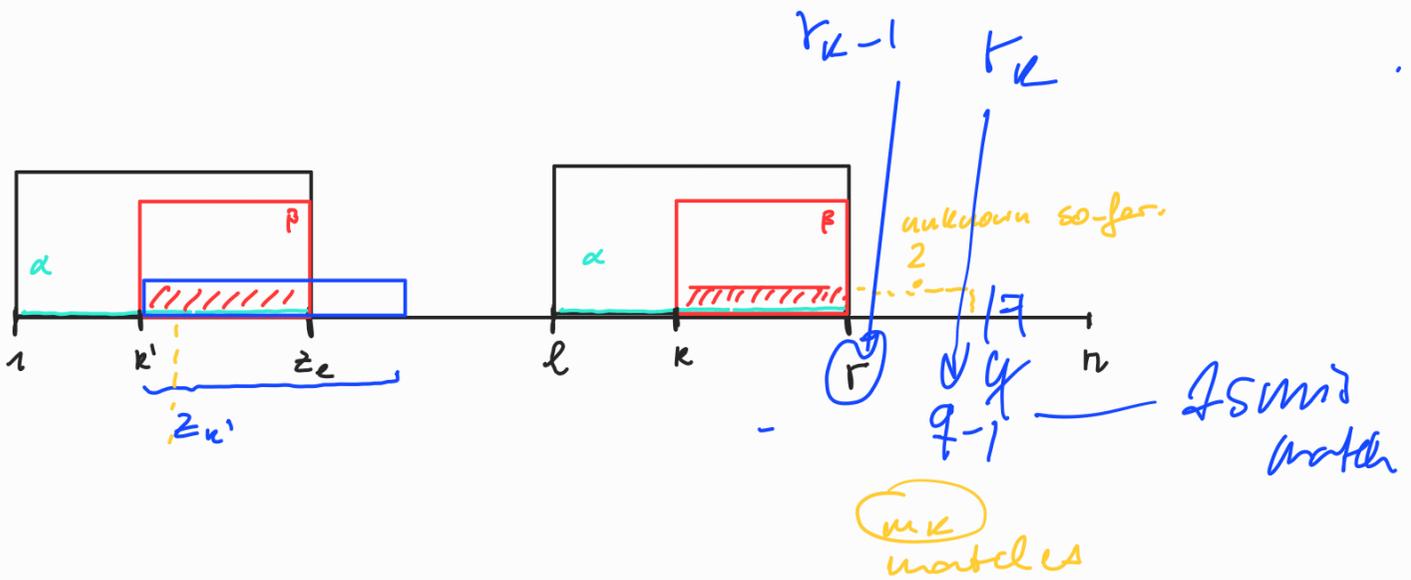
CASE 2A: in Alg  $r = r_{k-1}$  remains unchanged,

i.e.  $r_k = r_{k-1}$

moreover no comparisons are made, i.e.,  $m_k = 0$

$\Leftrightarrow r_k - r_{k-1} = m_k \checkmark$

CASE 2B: in Alg:  $q > r_{k-1}$  &  $r_k = q - 1$ :



$\Rightarrow r_k - r_{k-1} \geq m_k \checkmark$  [see picture].

$$n := |S| \Rightarrow \sum_{k=2}^n c_k \leq \sum_{k=2}^n 1 + \sum_{k=2}^n m_k = n - 1 + \sum_{k=2}^n m_k$$

(miss.)                      (match)

$\leq n - 1 + (r_2 - r_1) + (r_3 - r_2) + (r_4 - r_3) \dots + (r_n - r_{n-1}) = n - 1 + \underbrace{r_n - r_1}_{\leq n} \leq 2n - 1 \in O(n) \quad \square$

(claim) ↓  
in Alg  $r_1 = r_n = 0$

## Exmpl

i	1	2	3	4	5	6	7	8	9	10
S	a	c	a	c	a	b	a	c	a	c
$z_i$	-									
$l_i$	-									
$r_i$	-									

init  $r = l = 0$

$k=2$ :  $k=2 > r=0$  (Case 1)

compare  $S[2...n]$  with  $S[1..n]$

$\Rightarrow S(2) \neq S(1)$  mismatch

$\Rightarrow z_k = 0$ ,  $r, l$  unchanged.

i	1	2	3	4	5	6	7	8	9	10
S	a	c	a	c	a	b	a	c	a	c
$z_i$	-	0								
$l_i$	-	0								
$r_i$	-	0								

$k=3$ ,  $k=3 > r=0$  (Case 1)

compare  $S[3...n]$  with  $S[1..n]$

$S(3) = S(1)$

$S(4) = S(2)$

$S(5) = S(3)$

$S(6) \neq S(4)$

$\Rightarrow$

$z_k = 3 > 0$
$r = 3 + 3 - 1 = 5$
$l = 3$

$i$	1	2	3	4	5	6	7	8	9	10
$S$	a	c	a	c	a	b	a	c	a	c
$z_i$	-	0	3							
$l_i$	-	0	3							
$r_i$	-	0	5							

$$k=4, \quad r=5 \geq k=4 \quad (\text{Case 2})$$

$$(\alpha = aca, \beta = ca)$$

$$k^1 = k - l + 1 = 4 - 3 + 1 = 2$$

$$z_{k^1} = z_2 = 0 < r - k + 1 = 5 - 4 + 1 = 0$$

$$\Rightarrow \text{Case 2A: } z_4 = z_{k^1} = 0, \quad r, l \text{ unchanged.}$$

$i$	1	2	3	4	5	6	7	8	9	10
$S$	a	c	a	c	a	b	a	c	a	c
$z_i$	-	0	3	0						
$l_i$	-	0	3	3						
$r_i$	-	0	5	5						

$$k=5, \quad r=5 \geq k \quad (\text{Case 2 with } \alpha = aca, \beta = a)$$

$$k^1 = k - l + 1 = 5 - 3 + 1 = 3$$

$$z_{k^1} = z_3 = 3 > r - k + 1 = 5 - 5 + 1 = 1$$

$$\Rightarrow \text{Case 2A again and so on...}$$

$i$	1	2	3	4	5	6	7	8	9	10
$S$	a	c	a	c	a	b	a	c	a	c
$z_i$	-	0	3	0	1	0	4	0	2	0
$l_i$	-	0	3	3	5	5	7	7	9	9
$r_i$	-	0	5	5	5	5	10	10	10	10

Based on preprocessing with z-alg. we can design a:

## Simple linear-time exact matching Alg

Simple-exact-matching ( $P, T, \$$ ) // \$ character not in  $T \cup P$

- 1 occurrences =  $\emptyset$
- 2  $n = |P|, m = |T|$
- 3  $S = P\$T$
- 4 preprocess  $S$  with z-Alg to compute  $z_2 \dots z_{|S|}$
- 5 FOR ( $i = n+2, \dots, |S| = m+n+1$ )
- 6     IF ( $z_i = n$ )
- 7     └ add  $i-n-1$  to occurrences

Exmpl

$P = bbac, T = abba bbaca$

$i$	1	2	3	4	5	6	7	8	9	10	11	12	13	14
$S$	b	b	a	c	\$	a	b	b	a	<u>bb</u>	<u>aca</u>			
$z_i$						(doesn't matter...)	0	3	1	0	<span style="border: 1px solid red; padding: 2px;">4</span>	1	0	0

$z_{10} = |P| = 4 \Rightarrow P$  occurs on pos  
 $10 - 4 - 1 = 5$  of  $S$

Runtime:

Line 1	constant	
2	$n + m$	$\in O(n + m)$
3	$n + m + 1$	$\in O(n + m)$
4	$O( S )$	$= O(n + m)$
5-7	$m + n - 1 - (n + 2)$	$= O(m)$

TOTAL:  $O(n + m)$  time.

correctness:

Since  $\$$  not in  $P$  and  $T$   
 $\Rightarrow z_i \in |P| \forall i$

if  $|z_i| = |P| \Rightarrow$   $S[1 \dots |P|] \stackrel{\text{"pat } P}{=} S[i \dots i + n - 1] \stackrel{\text{"substring of } T}{=}$   
 $\Rightarrow P$  in  $S$  at pos  $i$   
 $= P$  in  $T$  at pos  $n - i - 1$

if  $P$  occurs in  $T$  at pos  $j$ , then  
 $P$  occurs in  $S$  at pos  $i = j + n + 1$   
 where  $i \in [n + 2, m + 2]$   
 $\Rightarrow |z_i| \geq |P|$

Since  $|z_i| \leq |P| \rightarrow |z_i| = |P|$  &  
 occurrence at pos  $j$   
 is reported

□

Thm: Simple-exact matching correctly reports  
 all occurrences of pattern  $P$  in text  $T$   
 in  $O(|P| + |T|)$  time.

[Back to slides]

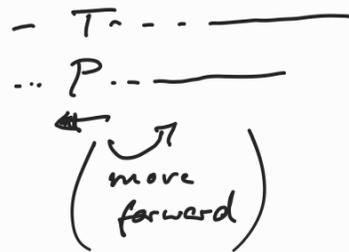
There is another important algorithm for pattern matching:

Boyer-Moore Algorithm is standard alg.

that is used in many applications for text search  
(e.g. google..)

BM-Alg. is based on 3 essential ideas:

- (1) Right-to-left scan
- (2) Bad-Character Rule
- (3) Good-Suffix Rule.



[ due to limited time & since we want to cover further topics, we skip this alg.

more information about this alg in any Brief-textbook. ]

So far: z - alg used to preprocess P & then find P.

Now: preprocess T instead!  
(text of static & remains unchanged (e.g. DNA))

## 8. Suffix Trees

classical "real world" problem:

For given Text & Pattern  $P$

where and how often does  $P$  in Text occur?

- Application:
- word processing
  - internet search
  - Bioinformatics ( $T = \text{genom}$ ,  $P = \text{gene sequ.}$ )
  - `fgrep` in Unix
  - search for plagiarism
  -

- Def:
- $\Sigma$  = finite alphabet,
  - string  $S$  (over  $\Sigma$ ) = sequence of characters in  $\Sigma$

Let  $S = x_1 x_2 \dots x_l$ ,  $x_i \in \Sigma$ ,  $1 \leq i \leq l$

$|S| = l$  = length of  $S$

$S[i..j]$  = substring  $x_i x_{i+1} \dots x_j$  of  $S$ , starting at pos  $i$   
ending at pos  $j$

[if  $i > j$ , then  $S[i..j] = \epsilon$  empty string]

$S[1..j]$  = prefix of  $S$

$S[i..l]$  = suffix of  $S$

prefix/suffix proper if it is not  $S$  or  $\epsilon$ .

$S(i)$  =  $i$ -th character of  $S$

$x, y \in \Sigma$  match if  $x = y$ , else mismatch

$\Sigma^*$  = set of all strings over  $\Sigma$ ,  $\Sigma^l$  = set of all strings over  $\Sigma$  of length  $l$

$S'$  substring of  $S$ , if  $\exists i, j$  st  $S' = S[i..j]$

Aim: given string  $P$  check where/how often  
 $P$  occurs in string  $T$

tree data structures can be seen as a collection of entities (= vertices) that are linked to simulate a hierarchy  
 = rooted tree

= trees  $T$  where one vertex  $f \in V(T)$  is called root

Given  $f \in V(T)$  we obtain a partial order  $\leq_T$  on  $V(T)$  st

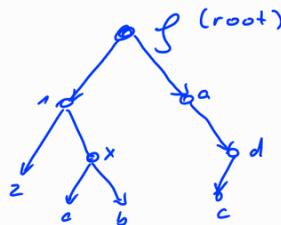
$x \leq_T y$  if  $y$  lies on simple path from  $f$  to  $x$ .

in this case:  $x$  is descendant of  $y$  &  $y$  is ancestor of  $x$

$v \in V(T)$  is common ancestor of vertices in  $W \subseteq V(T)$ , if  $w \leq_T v \forall w \in W$

a  $\leq_T$ -minimal common ancestor  $v$  of vertices in  $W$  is called least common ancestor ( $\text{lca}(W)$ )

Exmpl:



the arrows indicate  $\leq_T$

Note:  $x \leq_T x \forall x \in V(T)$

Notation:  $x \leq_T y$  if  $x \leq_T y$  &  $x \neq y$

Exmpl

f

1 has children 2 and x which are siblings

leaves are 2, a, b, c

root

children of  $x$ :  $y \in V(T)$  st  $y <_T x$  &  $(xy) \in E(T)$

$x$  is parent of its children

Siblings are vertices with the same parent

leaf: vertices with no children

Exmpl:  $S = \text{honolulu}$

$S[1..3] = \text{hon}$ , prefix & substring of  $S$ , no suffix

$S[5..8] = \text{lulu}$ , suffix & substring of  $S$ , no prefix

$S[5..7] = \text{lul}$ , substring of  $S$ , no prefix, no suffix

NOTE: empty string  $\epsilon$  is substring, prefix, suffix of all strings.

Naive way:



check occurrence of  $P$  in  $T$

by comparing  $1$  letter of  $P$  with  $i$  letter of  $T$   
 $j$  letter of  $P$  with  $i+j-1$  letter of  $T$   
 $n$  letter of  $P$  with  $i+n-1$  letter of  $T$

$\forall i = 1 \dots n-m+1$

$\Rightarrow$  run time  $O(n \cdot n)$

Often, the text is fixed & does not change  
(= long string)

- eg. • collected work of Shakespeare
- Genom

To find a pattern  $P$ , we need a datastructure that represents the text  
(= string) so that we can find efficiently  $P$

To this end: suffix trees

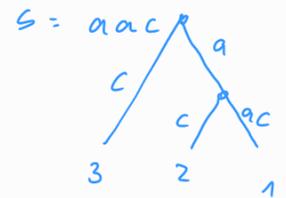
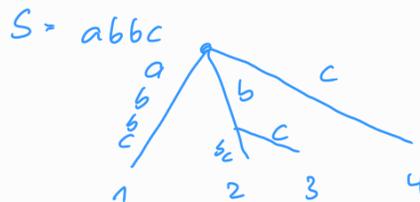
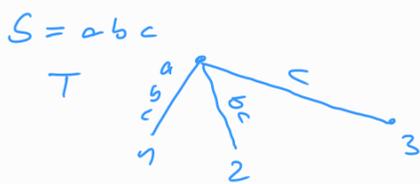
Def [suffix tree] let  $S$  be string of length  $|S|=m$

A suffix tree for  $S$  is a rooted tree  $T$  (with root  $r$ ) that satisfies the following properties:

- (X1)  $T$  has precisely  $m$  leaves that are uniquely labeled with  $1, 2, \dots, m$
- (X2) all inner vertices, (= vertices that are not leaves) except possibly the root, have at least 2 children
- (X3) every edge of  $T$  is labeled by a non-empty substring of  $S$
- (X4) For all distinct children  $v_1, v_2$  of  $v$  we have: label of edge  $(v, v_1)$  &  $(v, v_2)$  start with different characters
- (X5) if we concatenate the labels on the edges in order from  $r$  to leaf  $i$ , we obtain the suffix  $S[i..m]$

Exmpl:  $S = a$ ,  $T$  

[this is the only case where  $r$  has only one child, since if  $|S| > 1$  &  $T: [S[1..1]] \neq \epsilon$  we don't get  $S[m..m]$ , since  $[S[m..m]] (= 1)$  & (X3).]



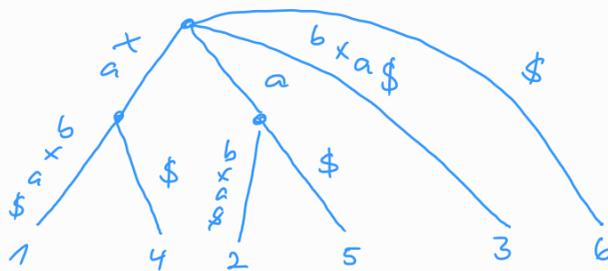


$\Rightarrow$  if suffix of  $S$  is also prefix of  $S$  no suffix tree exists!

IDEA: simply add special symbol  $\$$  at end of  $S$ ,  
 where  $\$$  does not occur in  $S$ .

From here on,  $S = x_1 \dots x_m$  with  $x_m = \$$

Example:  $S = xabxa\$$  (for "xabxa" no suffix tree)  
<sub>1 2 3 4 5 6</sub>



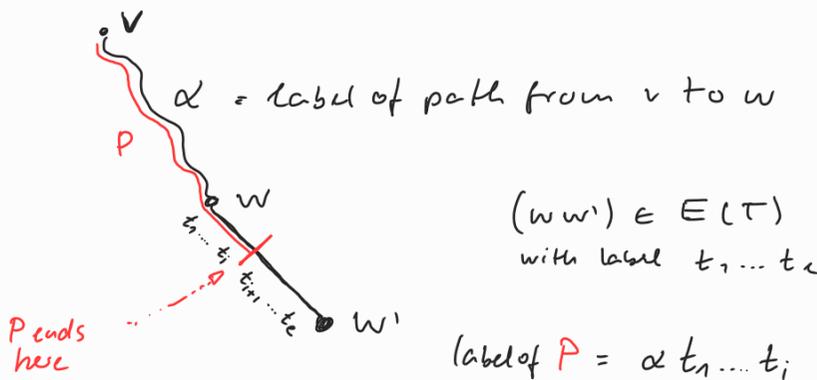
Def.

Let  $v \leq_T w$  &  $P$  be the unique simple path from  $v$  to  $w$



label of  $P$  = concatenation of labels of edges  
 in order from  $v$  to  $w$ .

By slight abuse of notation, we also consider simple paths  
 in suffix tree that may end on edge (and not on vertex)



# How to construct suffix tree & how does this help?

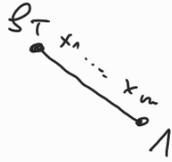
## Alg. SUFFIXTREE (Idea)

// iteratively build suffix trees

$T_1 \dots T_m$  by adding  $S[1..m], \dots, S[m..m]$   
in previously constructed  $T_i$

Input:  $S = x_1 \dots x_m$

1) Construct  $T_1 =$



2) Assume  $T_i$  is constructed, then construct  $T_{i+1}$   
( $i < m$ )  
as follows:

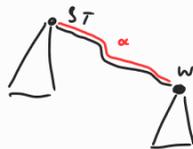
(a) Find path  $P$  in  $T_i$  that starts in  $ST_i$   
with longest label that is a prefix of  
 $S[i+1..m]$  [path could end in  $\{\}$ ]

(i) By similar arguments as in proof of L. 8.1  
this path is uniquely determined  
since no two edges  $(v, v_1), (v, v_2)$ ,  $v_1, v_2$  children of  $v$   
have labels starting with same symbol ( $\neq \emptyset$ ).

(ii) this path will never end in leaf of  $T_i$   
since  $P$  starts at root & concatenation of edge labels  
to leaf  $i$  yields  $S[i..m]$  where  $|S[i..m]| > |S[i+1..m]|$   
[by induction - EXERC]

(b) This path  $P$  either ends in edge or non-leaf vertex of  $T_i$

$P$  ends in vertex:

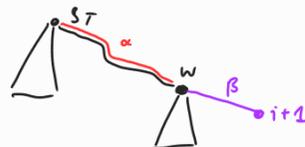


$w$  no leaf  $\Rightarrow w$  has children.

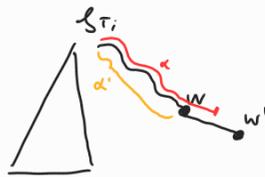
$\alpha$  prefix of  $S[i+1..m]$ ,  $m = j$   
But  $\alpha \neq S[i+1..m]$   
otherwise  $S[i+1..m] = S[l..m]$   
for some  $l$  but  $l < i+1$

$\Rightarrow \alpha = S[i+1..m-j]$ ,  $j > 1$

$\Rightarrow$  add edge  $(w, i+1)$   
with label  $\beta = S[m-j+1..m]$



P ends in edge:



$$\alpha = S[i+1 \dots m-j]$$

Label edge  $(w, w') = t_1 \dots t_r t_{r+1} \dots t_s$

$$\Rightarrow \alpha = \alpha' t_1 \dots t_r \text{ for some } 1 \leq r < s$$

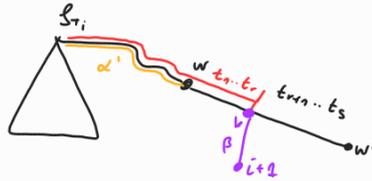
add new vertex  $v$   
on edge  $(w, w')$

& put  $\text{label}(wv) = t_1 \dots t_r$

$$\text{label}(vw') = t_{r+1} \dots t_s$$

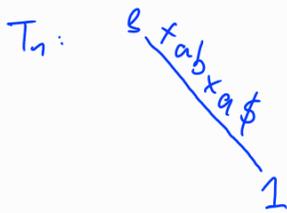
+ add new edge  $(v, i+2)$

$$\text{with label}(v, i+2) = S[m-j+2 \dots m] = \beta$$

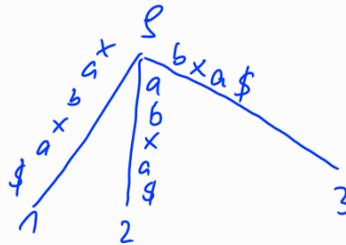


For both cases,  $T_{i+2}$  contains now path from  $s_{i+1}$  to  $i+2$   
with label  $S[i+2 \dots m]$

Exmpl:  $S = xabxa\$$

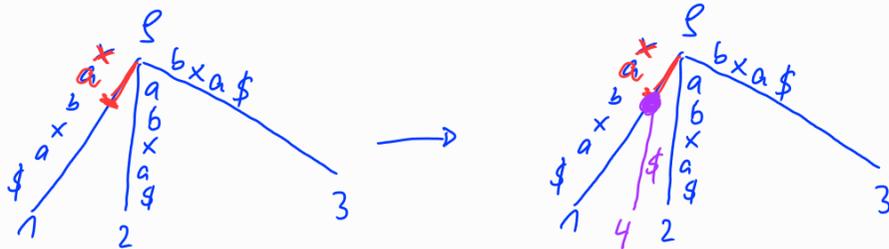


$T_2/T_3$  (longest path ends in  $\$$ )

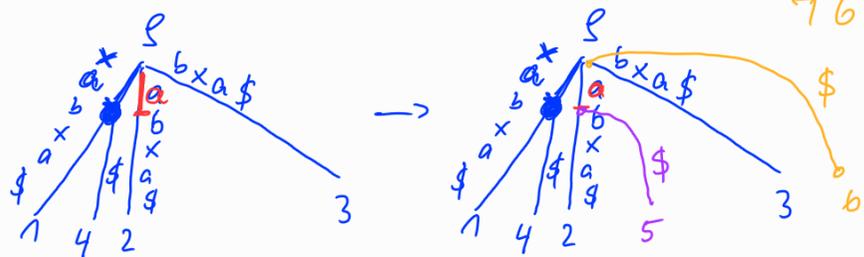


$T_4$ :  $S[4..m] = xa\$$

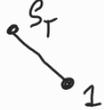
longest path P



$T_5$ :  $S[5..m] = a\$$



# SUFFIX-TREE(S)

init  $\mathcal{T}$  as  and label  $(s_{\mathcal{T}}, 1) = S[1..m]$

FOR (  $i=2..m$  ) DO

(end,  $i'$ ,  $l'$ ) = FIND-LONGEST-PATH ( $\mathcal{T}$ ,  $S[i..m]$ )

returns end of path, that is,  
 either end = v or end = e  
 & index  $i'$  &  $l'$

IF (end = edge e) //  $e = (uv)$  with label  $\gamma$

remove  $e = (uv)$   
 add new vertex  $v$   
 add new edges  $e_1 = (uv)$ ,  $e_2 = (v, w)$   
 & label  $e_1 = \gamma[1..l']$   
 label  $e_2 = \gamma[l'+1..|\gamma|]$



add edge  $(v, i)$  to  $\mathcal{T}$  with label  $S[i+i'..m]$

## FIND-LONGEST-PATH ( $\mathcal{T}$ , $s'$ )

[Sketch  $\rightarrow$  more details in Appendix]

"Follow path from  $s_{\mathcal{T}}$  to  $v/c$  as long as possible, i.e., as long as letters on this path match with letters  $s' = x_{i'}..x_m$ "

& return corresponding positions  $i'$ ,  $l'$

+ if end is edge or vertex.

## Theorem [Ukkonen]:

[without proof]

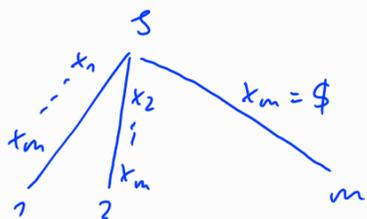
for  $S$  of length  $m$  the suffix tree can be constructed in  $O(m)$  time

[quite sophisticated pointer-adjustments].

# Space complexity

Suffix tree without labels  $O(m)$   
but we need to store labels!

worst case,



each edge label of  $(\ell, i)$  is of size  $i$   
 $\Rightarrow \sum_{i=1}^m i = O(m^2)$  space

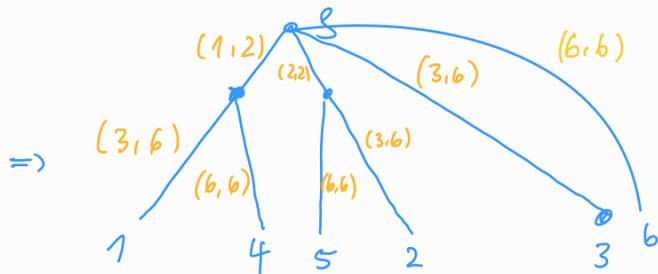
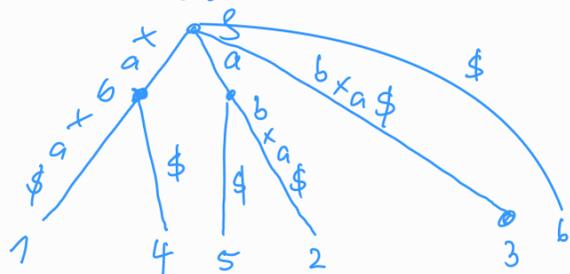
$O(m^2)$  space is bad (eg. genome of small bacteria has  $10^6$  characters  $\sim$  1TB storage)

IDEA:

instead of saving label  $S[i..j]$  for  $c$   
we only save pair  $(i, j)$   
= compressed suffix tree

$\Rightarrow O(m)$  space (same space as text  $S$ )

Exmpl:  $S = xabxa\$$   
1 2 3 4 5 6



Why Suffix trees?

A: Examples!

In what follows, we assume to have Ukkonen's version:  $O(m)$  time

Exmpl 1

"exact text-search"

Given (long) text  $S$  & pattern  $P$  (=string)

Does  $P$  in  $S$  occur?

Let  $T$  be suffix tree of  $S$ .

Call  $FLP(T, P)$  once  $O(|P|)$  time

return value is (end, index  $i$ , ...)

st  $P[1..i']$  corresponds to label of path in  $T$  that starts in  $\mathcal{L}T$

Observe, any path in  $T$  from  $\mathcal{L}T$  to leaf  $i$  corresponds to  $S[i..m]$

&  $P[1..i']$  corresponds therefore to a prefix of some suffix of  $S$ , that is, a substring of  $S$ .

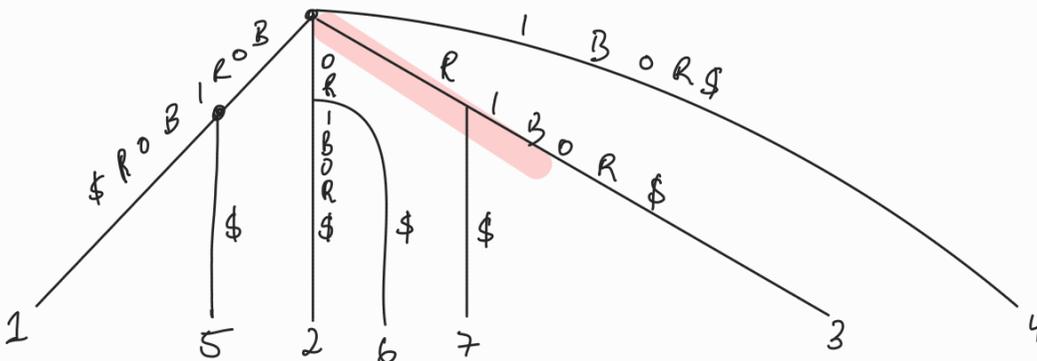
$\Rightarrow$  if  $i' = |P| \Rightarrow P$  occurs in  $T$  else, not.

$\Rightarrow$  runtime  $O(|P|)$

Exmpl:

$T = \text{\$ORIBOR\$}$

$P = \text{RIB}$



[every exist letter appears once at first letter of edge ( $\mathcal{L}V$ ) for some  $v$ .]

## Exmpl 2 "substring - database - search"

Given strings  $S_1 \dots S_e$  (Database of Texts)

Does  $P$  occur in at least one of the  $S_i$ ?

$\Rightarrow$  let  $\$1 \dots \$e$  pairwise distinct symbols that do not occur in any  $S_i$  &  $P$

put  $S = S_1 \$1 S_2 \$2 \dots S_e \$e$

Construct suffix tree  $\mathcal{T}$  for  $S$  ( $O(|S_1| + |S_2| + \dots + |S_e|)$  time)

use IDEA of Exmpl 1

## Exmpl 3

### Find all occurrences of $P$

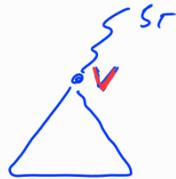
given  $S$  with suffix tree  $\mathcal{T}$  & pattern  $P$  with  $|P|=n$

Find all occurrences of  $P$  in  $\mathcal{T}$ , that is,

all indices  $i$  st  $P = S[i:i+n-1]$

Call  $FLP(\mathcal{T}, P)$

$\rightarrow$  this gives  $end = v$  or  $end = edge(uv)$

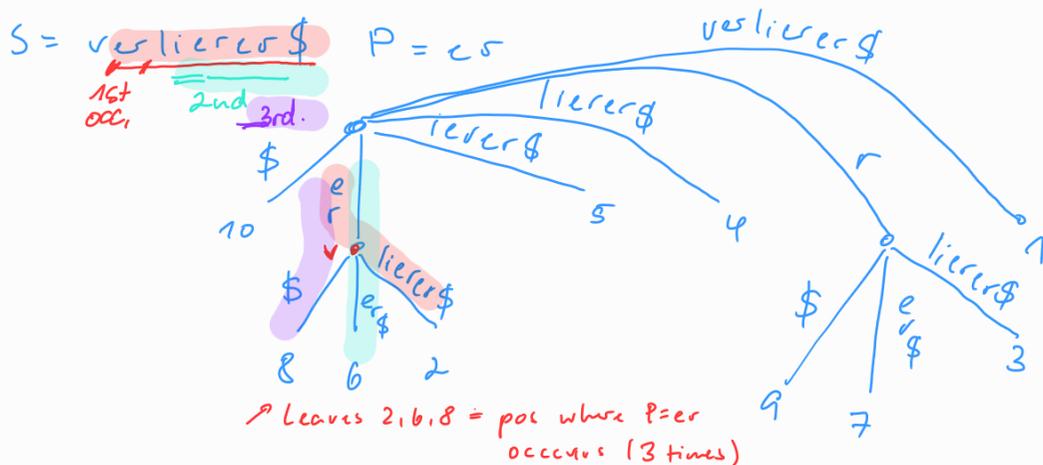


# leaves  $i \leq v = \#$  occurrences of  $P$  in  $S$

&  $P$  occurs in pos  $i$   $\forall$  all leaves  $i \leq v$ .

[Exercise: works in  $O(|P| + k)$  time]

$k = \#$  of occurrences of  $P$



Exmpl 4

Find longest substring in  $S$  that occurs at least 2 times

[Application: Find repeated regions in genome]

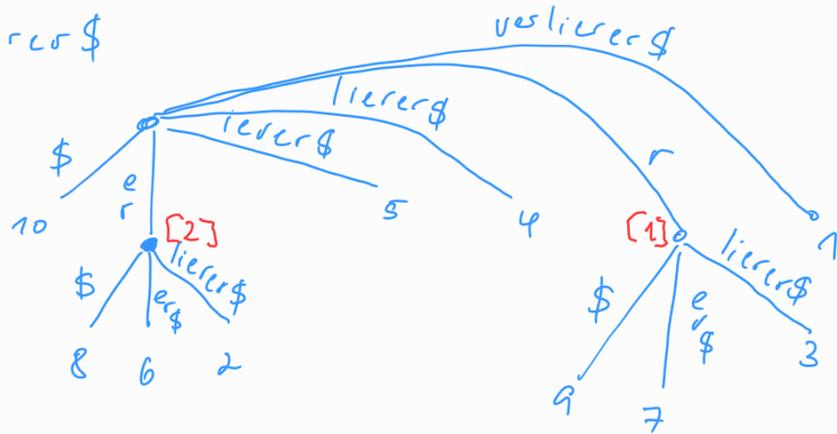
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string depth of  $v$  in  $T$  is  $|a|$



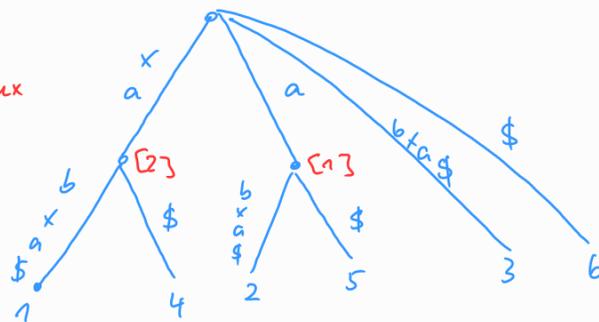
$S = \text{verlierer}\$$

string depth of inner vertex



$S = \text{xabxa}\$$

string depth of inner vertex



$\text{x}a$  occurs 2 times.

- every inner vertex has 2 children & first symbol on edges with same parent have distinct symbols (at least)

$\Rightarrow$  length of longest substring that occurs at least 2 times = maximum string-depth of inner vertices.

[Exus.: works in  $O(|S|)$  time.]

# Exmpl 5

## efficient detection of pairwise overlaps

Given  $J = \{s_1 \dots s_N\}$  set of strings (eg. sequenced DNA fragments)

Aim: define  $ov(s_i, s_j) \neq r_{ij}$   
"size of largest overlap"

$$S = s_1 \$1 s_2 \$2 \dots s_N \$N$$



$$s_1 = ABA$$

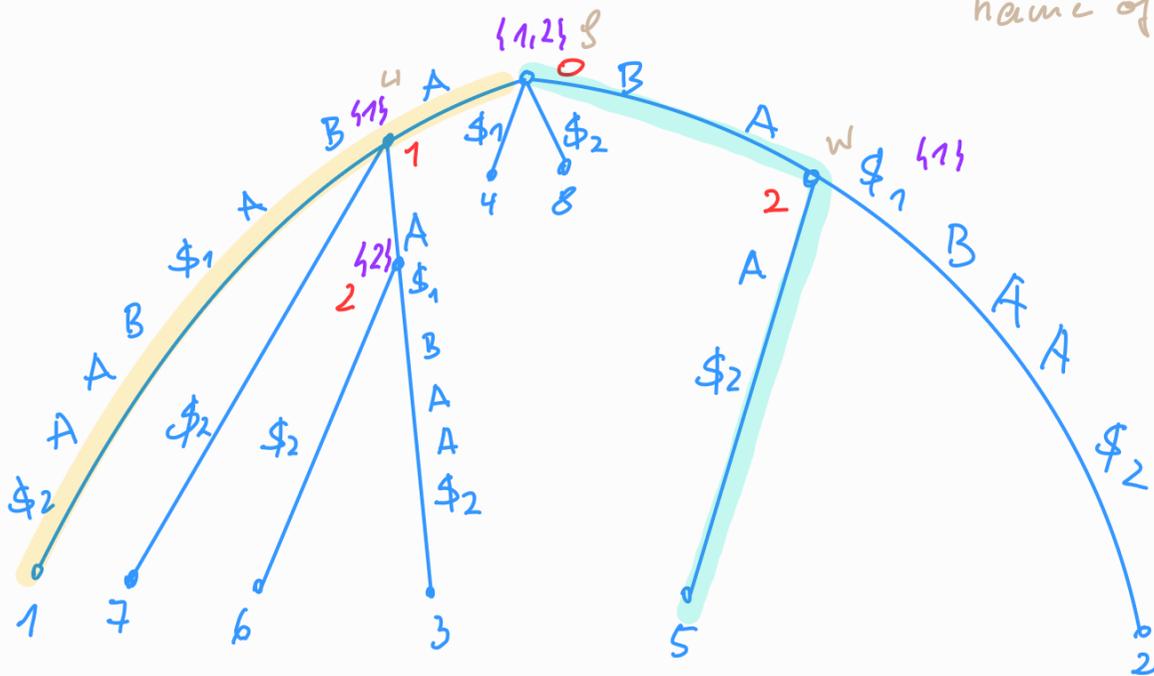
$$\rightarrow ov(s_1, s_2) = 2$$

$$s_2 = BAA$$

$$ov(s_2, s_1) = 1$$

$$S = \overset{1}{A} \overset{2}{B} \overset{3}{A} \$1 \overset{4}{B} \overset{5}{A} \overset{6}{A} \$2$$

name of vertex



$P_1 =$  path starting with label  $s_1 \$1$

$P_2 =$  path starting with label  $s_2 \$2$

for non-leaf  $x$ :  $L(x) = \{i \mid \exists \text{ leaf } v \text{ st } (xv) \text{ edge in suffix tree} \& \text{ label } (xv) \text{ starts with } \$i\}$ .

string depth of  $x =$  # characters on path from root to  $x$ .

For ( $j=1 \dots N$ )

```
x = f (root)
WHILE (x not leaf)
  FOR (all  $i \in L(x), i \neq j$ )
    IF ( $\text{depth}(x) < \min(|S_i|, |S_j|)$ 
        OR  $(S_i, S_j) = \text{depth}(x)$ )
      x ← child of x on  $\tau_j$ 
```

$j=1, x=f$

FOR ( $i \in L(f) = \{1, 2\}, i \neq 1$ )

$\Rightarrow i=2$

$\text{depth}(f) = 0 < \min(|S_1|, |S_2|)$

$\Rightarrow \underline{\text{OR}(S_2, S_1) = 0}$

$x \leftarrow u$

FOR ( $i \in L(u) = \{1\}, i \neq 1$ )

not any

$x \leftarrow \text{leaf stop.}$

$j=2, x=f$

FOR ( $i \in L(f) = \{1, 2\}, i \neq 2$ )

$\Rightarrow i=1$

$\text{depth}(f) = 0 < \min(|S_1|, |S_2|)$

$\text{OR}(S_1, S_2) = 0$

$x \leftarrow w$

FOR ( $i \in L(w) = \{1\}, i \neq 2$ )

$\Rightarrow i=1$

$\text{depth}(w) = 2 < \min(|S_1|, |S_2|)$

$\Rightarrow \underline{\text{OR}(S_1, S_2) = 2}$

$x \leftarrow \text{leaf stop.}$

[without proof of correctness or runtime analysis]

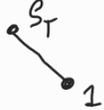
$O(N \|S\|)$

Exercise  $\rightarrow$  Book "Alg. Aspects of Binding."

## Appendix

### Details of $O(n^2)$ Algorithm

# SUFFIX-TREE(S)

init  $\mathcal{T}$  as  and label  $(s_{\mathcal{T}}, 1) = S[1..m]$

FOR (  $i=2..m$  ) DO

(end,  $i'$ ,  $l'$ ) = FIND-LONGEST-PATH ( $\mathcal{T}$ ,  $S[i..m]$ )

// returns end of path, that is,  
// either end = v or end = e  
// & index  $i'$  &  $l'$

IF (end = edge e) //  $e = (uw)$  with label  $\gamma$

remove  $e = (uw)$   
add new vertex  $v$

add new edges  $e_1 = (uv)$ ,  $e_2 = (v, w)$   
& label  $e_1 = \gamma[1..l']$   
label  $e_2 = \gamma[l'+1..|\gamma|]$



add edge  $(v, i)$  to  $\mathcal{T}$  with label  $S[i+i'..m]$

## FIND-LONGEST-PATH ( $\mathcal{T}$ , $S'$ )

$j=1$ ,  $v = s_{\mathcal{T}}$

WHILE ( $j \leq |S'|$ ) DO

Find edge  $e = (vw)$  in  $\mathcal{T}$ ,  $w \in \mathcal{T}_v$ , whose label starts with  $S'(j)$

IF (such edge does not exist)

└ RETURN ( $v, j-1, \emptyset$ ) // path ends in  $v$

Let  $\gamma$  be label of  $e$

$l = 1$

WHILE ( $j \leq |S'|$  &  $l \leq |\gamma|$  &  $S'(j) = \gamma(l)$ ) DO

└  $j = j+1$   
 $l = l+1$

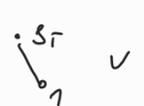
IF ( $l \leq |\gamma|$ ) // while loop ended at some point before " $S'(j) = \gamma(l)$   
//  $\forall l = 1..|\gamma|$ "  
└ RETURN ( $e, j-1, l-1$ )  
ie,  $S(j) \neq S(e)$

$v = w$  // go to consider edges starting at  $w$

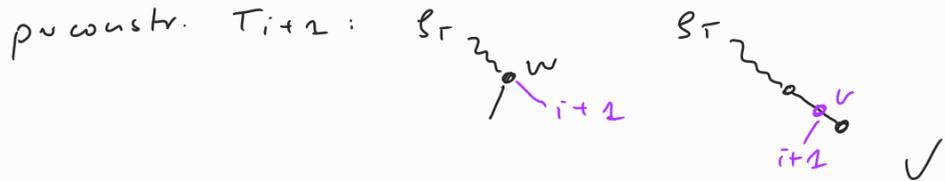
RETURN ( $v, |S'|, \emptyset$ )

Prop. 8.2

Alg. SUFFIXTREE correctly computes suffix tree for  $S = x_1 \dots x_m$ ,  $x_m = \$$ .

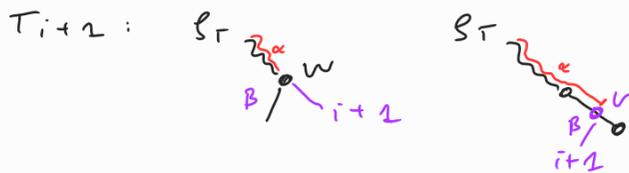
proof: (X1):  $T_1$  has 1 leaf,  $T_2$  has 2 leaves ...  $T_m$  has  $m$  leaves ✓  
 (sketch) (X2):  $T_1$ : 

Assume, for  $T_i$ , each inner vertex (except possibly  $f$ ) has at least 2 children



(X3) by construction & induction each edge has non-empty string as label

(X4)  $T_i \cup$  Assume for  $T_i \forall v$ : label  $(v_1), (v_2)$  starts with different symbol, where  $v_1, v_2$  are children of  $v$ .



Since  $\alpha$  is longest substring that is prefix of

$$S[i+1..m] = \underbrace{x_{i+1} \dots x_r}_{\alpha} \dots \underbrace{x_{r+1} \dots x_m}_{\beta}$$

$\beta(1) \neq x_{r+1}$  as otherwise a not longest prefix! ✓

(X5) by const. & induct. concatenating edge-labels from  $f$  to  $i$  yields  $S[i..m]$ .

□

## Runtime:

SUFFIXTREE(S),  $S = x_1 \dots x_m$

add/remove etc in  $O(1)$  time

→  $m$  times FIND-LONGEST-PATH (FLP)  
it called.

FLP: • in each of the  $|S|$ -steps in 1st while-loop

Find-edge → for  $v$  all neighbors  $w$  are considered

⇒  $\deg_{\rightarrow}(v)$  many vertices

over all calls of 1st-while loop, "Find-edge" is called,

$$\leq \sum_{v \in V} \deg(v) = 2|E| = O(|E|) \text{ times} \\ = O(m)$$

• 1st + 2nd while loop:

WHILE ( $j \in |S|$ ) DO

⋮

WHILE ( $j \in |S|$  &  $l \in |x|$  &  $S'(j) = x(l)$ ) DO

$j = j + 1$   
    ⋮

⇒  $j \leq m \Rightarrow O(m)$  time

• Remaining operation in 1st-while loop:  $O(1)$

⇒ FLP runs in  $O(m)$  time

⇒ SUFFIX-TREE(S) runs in  $O(m^2)$  time.

Theorem [Ukkonen]: for  $S$  of length  $m$  the suffix tree can  
[without proof] be constructed in  $O(m)$  time