

## String Matching and Suffix Tree

Gusfield Ch1-7

EECS 458  
CWRU  
Fall 2004

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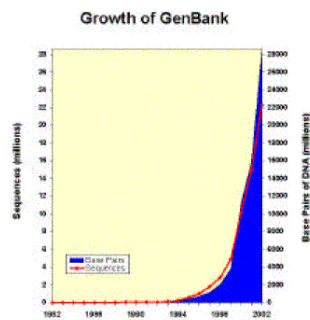
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## GenBank



Size doubles every 14 months.  
As of the end of 2002,  
GenBank contained  
approximately  
28,507,990,166 bases  
in 22,318,883 sequence  
records

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## BLAST (Altschul et al'90)

- Idea: true match alignments are very likely to contain a short segment that identical (or very high score).
- Consider every substring (seed) of length  $w$ , where  $w=12$  for DNA and 4 for protein.
- Find the exact occurrence of each substring (how?)
- Extend the hit in both directions and stop at the maximum score.

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## Problems

- Pattern matching: find the exact occurrences of a given pattern in a given structure ( string matching)
- Pattern recognition: recognizing approximate occurrences of a given pattern in a given structure (image recognition)
- Pattern discovery: identifying significant patterns in a given structure, when the patterns are unknown (promoter discovery)

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## Definitions

- String  $S[1..m]$
- Substring  $S[i..j]$
- Prefix  $S[1..i]$
- Suffix  $S[i..m]$
- Proper substring, prefix, suffix
- Exact matching problem: given a string  $P$  called pattern, and a long string  $T$  called text, find all the occurrences of  $P$  in  $T$ .

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## Naïve method

- Align the left end of  $P$  with the left end of  $T$
- Compare letters of  $P$  and  $T$  from left to right, until
  - Either a mismatch is found (not an occurrence)
  - Or  $P$  is exhausted (an occurrence)
- Shift  $P$  one position to the right
- Restart the comparison from the left end of  $P$
- Repeat the process till the right end of  $P$  shifts past the right end of  $T$
- Time complexity: worst case  $\theta(mn)$ , where  $m=|P|$  and  $n=|T|$
- Not good enough!

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- When mismatch occurs, shift P more than one letter, but never shift so far as to miss an occurrence
- After shifting, skip over parts of P to reduce comparisons
- Preprocessing of P or T

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- Can be on pattern  $P$  or text  $T$ .
- Given a string  $S$  ( $|S|=m$ ) and a position  $i > 1$ , define  $Z_i$ : the length of longest common prefix of  $S$  and  $S[i..m]$
- Example,  $S=abxyabxz$



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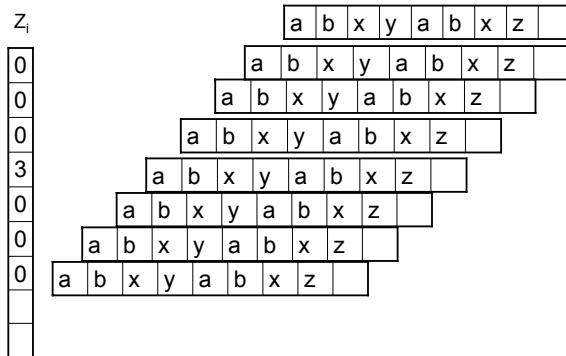
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## Fundamental preprocessing



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## Fundamental preprocessing

- Intention:
  - Concatenate P and T, inserted by an extra letter \$:  $S = P\$T$
  - Every  $i$ ,  $Z_i \leq |P|$
  - Every  $i > |P| + 1$  and  $Z_i = |P|$  records the occurrences of P in T
- Question: running time to compute all the Zs? The naïve method according to the definition runs in  $\theta((m+n)^2)$  time!

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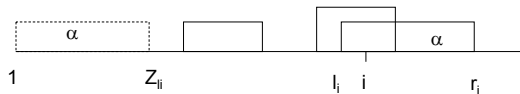
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## Fundamental preprocessing

- Goal: linear time to compute all the Zs
- Z-box: for  $Z_i > 0$ , it is the box starting at  $i$  with length  $Z_i$  (ending at  $i + Z_i - 1$ ),
- $r_i$ : the rightmost end of a  $Z_j$ -box ( $j + Z_j - 1$ ) for all  $1 < j \leq i$  such that  $Z_j > 1$ .
- $l_i$ : the left node of  $Z_j$ -box ending at  $r_i$




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## Fundamental preprocessing

- Computing  $Z_k$ :
  - Given  $Z_i$  for all  $1 < i < k$
  - Let  $r = r_{k-1}$  and  $l = l_{k-1}$
  - 1.  $k > r$ : compute  $Z_k$  explicitly (updating accordingly  $r$  and  $l$  if  $Z_k > 0$ )
  - 2.  $k \leq r$ :  $k$  is in the  $Z$ -box starting at  $l$  (substring  $S[l..r]$ ), therefore  $S[k] = S[k-l+1]$ ,  $S[k+1] = S[k-l+2]$ , ...,  $S[r] = S[Z_l]$ . In other words,  $Z_k \geq \min\{Z_{k-l+1}, r-k+1\}$




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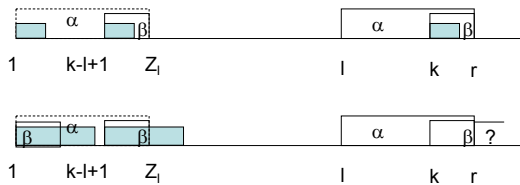
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## Fundamental preprocessing

- A)  $Z_{k-l+1} < (r-k+1)$ :  $Z_k = Z_{k-l+1}$ , and  $r, l$  remain unchanged
- B)  $Z_{k-l+1} \geq (r-k+1)$ :  $Z_k \geq (r-k+1)$  and start comparison between  $S[r+1]$  and  $S[r-k+2]$  until a mismatch is found (updating  $r$  and  $l$  accordingly if  $Z_k \geq r-k+1$ )



## Fundamental preprocessing

- Conclusions:
  - $Z_k$  is correctly computed
  - There are a constant number of operations besides comparisons for each  $k$ 
    - $|S|$  iterations
    - Whenever a mismatch occurs, the iteration terminates
    - Whenever a match occurs,  $r$  is increased
  - In total at most  $|S|$  mismatches and at most  $|S|$  matches
  - Running time  $\theta(|S|)$  and space  $\theta(|S|)$

## Fundamental preprocessing

- Th: there is a  $\theta(n+m)$ -time and space algorithm which finds all the occurrences of  $P$  in  $T$ , where  $m=|P|$  and  $n=|T|$ .
- Notes:
  - Alphabet-independent
  - Space requirement can be reduced to  $\theta(m)$
  - Not well suited for multiple patterns searching
  - Strictly linear, every letter in  $T$  has to be compared at least once

## Projects

- Topics
- Meeting: 3 times, as a group
- Presentations: 25 minutes/student (~20m talk + 5m questions)
- Term paper: single space, 11pt, 1in margin. 5-6p, 9-10p 10-12p, exclude references

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## The Boyer-Moore algorithm: an example

- $P = abxabxab$ ,  $T = daaabbxabxab$

|   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| d | a | a | a | b | x | a | b | a | b | x | a | b | x | a | b |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|

|   |   |   |   |   |   |   |   |
|---|---|---|---|---|---|---|---|
| a | b | x | a | b | x | a | b |
|---|---|---|---|---|---|---|---|

|   |   |   |   |   |   |   |   |
|---|---|---|---|---|---|---|---|
| a | b | x | a | b | x | a | b |
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|   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| d | a | a | a | b | x | a | b | a | b | x | a | b | x | a | b |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|

|   |   |   |   |   |   |   |   |
|---|---|---|---|---|---|---|---|
| a | b | x | a | b | x | a | b |
|---|---|---|---|---|---|---|---|

|   |   |   |   |   |   |   |   |
|---|---|---|---|---|---|---|---|
| a | b | x | a | b | x | a | b |
|---|---|---|---|---|---|---|---|

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## The Boyer-Moore algorithm

- Rule 1: right-to-left comparison
- Rule 2: Bad character rule
  - For each  $x \in \Sigma$ ,  $R(x)$  denotes the right-most occurrence of  $x$  in  $P$  (0 if doesn't appear)
  - When a mismatch occurs,  $T[k]$  against  $P[i]$ , shift  $P$  right by  $\max\{1, i - R(T[k])\}$  places. This takes  $T[k]$  against  $P[R(T[k])]$
  - $|\Sigma|$  space to store  $R$ -values
- Rule 3: good suffix rule

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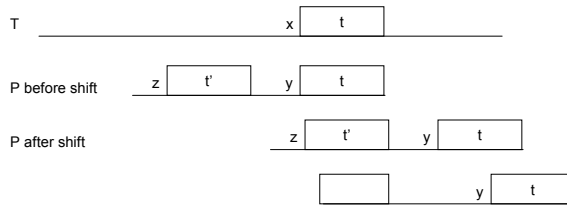
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## The Boyer-Moore algorithm




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## The Boyer-Moore algorithm

- Rule 3: good suffix rule
  - When a mismatch occurs,  $T[k]$  against  $P[i]$
  - Find the rightmost occurrence of  $P[(i+1)..m]$  in  $P$  such that the letter to the left differs  $P[i]$
  - Shift  $P$  right such that this occurrence of  $P[(i+1)..m]$  is against  $T[(k+1)..(m+k-i)]$
  - If there is no occurrence of  $P[(i+1)..m]$ , find the longest prefix of  $P$  matches a suffix of  $P[(i+1)..m]$ , shift  $P$  right such that this prefix is against the corresponding suffix

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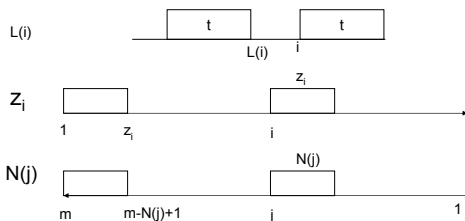
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## Preprocessing for the good suffix rule

- Let  $L(i)$  denote the largest position less than  $m$  such that string  $P[i..m]$  matches a suffix of  $P[1..L(i)]$
- Let  $N(j)$  denote the longest suffix of substring  $P[1..j]$  that is also a suffix of  $P$
- Recall  $Z$ , the length of longest substring of  $P$  starts 1 and matches a prefix of  $S$ .




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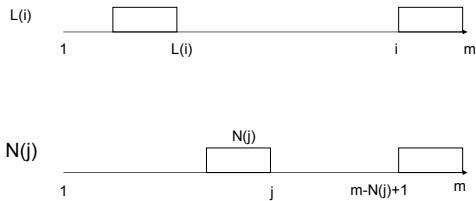
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## Preprocessing for the good suffix rule

- Thm:  $L(i)$  is the largest index  $j$  less than  $m$  such that  $N(j) \geq |P[i..m]| = m - i + 1$ .




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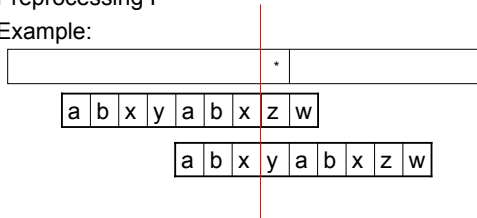
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## The Knuth-Morris-Pratt Algorithm

- History:
  - Best known
  - Not the method of choice, inferior in practice
  - Can be generalized for multiple string matching
- Preprocessing P
- Example:




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## KMP

- Idea:
  - left to right comparison,
  - Shift P more places without missing occurrence
- A prefix of P matches a proper suffix of  $P[1..i]$  and the next letters do not match!
- Define  $s_i$  of P,  $2 \leq i \leq m$  the length of longest proper suffix of  $P[1..i]$  that matches a prefix of P,  $s_1 = 0$ .

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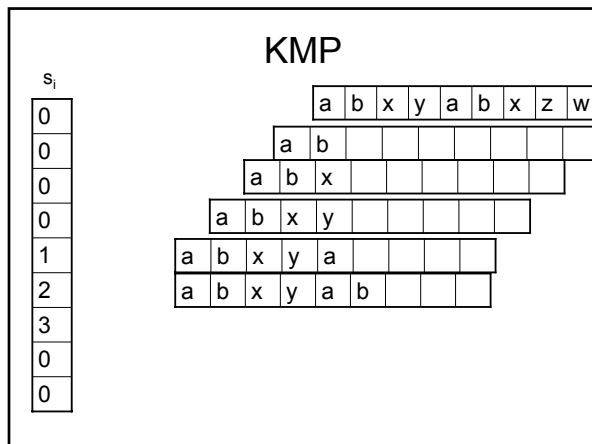
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### KMP

- Define  $s'_i$ ,  $2 \leq i \leq m$ , the length of the longest proper suffix of  $P[1..i]$  that matches a prefix of  $P$  with the additional condition that character  $P[i+1]$  differs from  $P[s'_i+1]$ .
- Obviously  $s'_i \leq s_i$  for any  $i$

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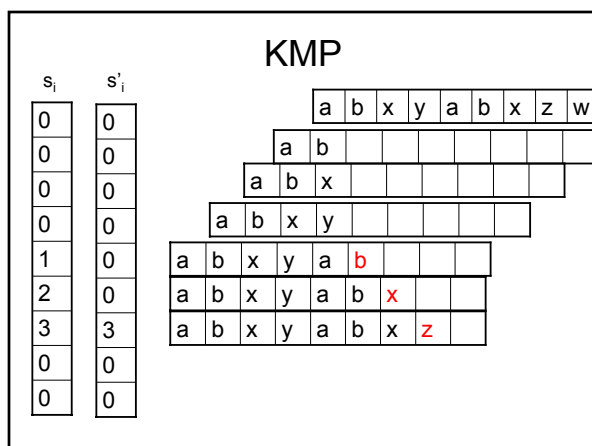
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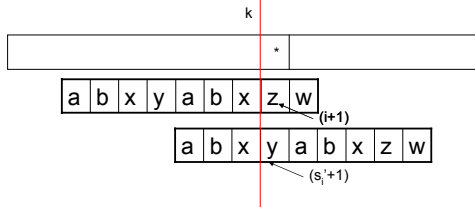
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## KMP

- Shifting rule: shift P to the right ( $i - s_i'$ ) spaces (  $i=n$  if an occurrence)



- Since  $P[1..s_i']$  matches  $P[i-s_i'+1, i]$  and  $P[i-s_i'+1, i]$  matches  $T[k-s_i'+1, k]$ , thus we skip  $s_i'$  comparisons.

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## KMP

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KMP(){
  m=|P|, n=|T|, s', q=0;
  for (i=0; i<n; i++){
    while q>0 and P[q+1] <> T[i]
      q=s_q'
    if(P[q+1]==T[i])
      q++;
    if(q==m) {
      find a pattern at position i-m, q=s_q'
    }
  }
}

```

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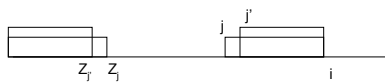
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## KMP

- Correctness
- $Z_j$ : the length of the longest common prefix of P and  $P[j..n]$
- $s_i'$  the length of the longest proper suffix of  $P[1..i]$  that matches a prefix of P, with the additional condition that the character  $P[i+1]$  differs from  $P[s_i'+1]$



- Therefore,  $s' = \max\{Z_j \mid Z_j = i - j + 1\}$

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## KMP

- Running time:
  - In total  $s$  phases (of comparison/shift),  $s \leq n$
  - Every 2 consecutive phases overlap one letter (the mismatched one) from  $T$
  - Therefore, in total  $n+s \leq 2n$
- Questions:
  - Any letter from  $T$  is skipped for comparison?
  - The times of comparison for a letter is at most a constant time? (real time algorithm)

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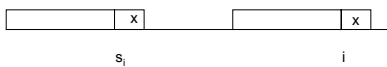
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## Original preprocessing of KMP: calculating $s'$ directly

- $s_i$ : the length of longest proper suffix of  $P[1..i]$  that matches a prefix of  $P$
- $s'_i$ , the length of the longest proper suffix of  $P[1..i]$  that matches a prefix of  $P$  and  $P[i+1] \neq P[s'_i+1]$ .
- So  $s'_i = s_i$  if  $P[i+1] \neq P[s_i+1]$   
 $s'_i = s'_{s_i}$  otherwise




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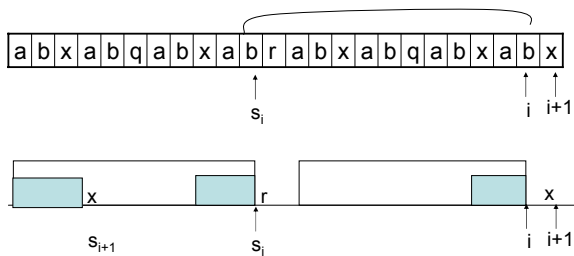
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## Calculate $s$

$$s_{i+1} = s_i + 1 \text{ if } P[s_i+1] = P[i+1]$$

$$= s_{(s_i)} + 1 \text{ if } P[s_i+1] \neq P[i+1]$$

$$= \dots$$




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[illegible]

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# Multiple pattern matching problem

- Given a set of pattern  $P = \{P_1, P_2, \dots, P_k\}$  and a text (databases)  $T$ , find all the occurrences of all the patterns...
- Keyword tree: given a set of pattern  $P = \{P_1, P_2, \dots, P_k\}$ , the keyword tree  $K$  is a tree:
  - Rooted, directed
  - Each edge is labeled with one letter
  - Edges coming out of a node have distinct labels
  - Every pattern is spelled out (map to one node)
  - Every leaf maps to some pattern

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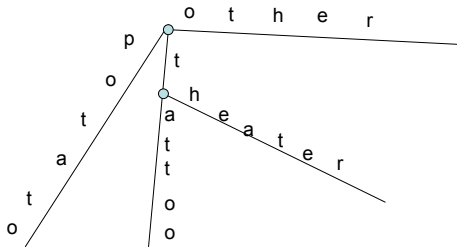
# An example

- abxabqabxabraxabqabxabx

- abc

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- $P = \{\text{potato, tattoo, theater, other}\}$



- Linear time construction
- Dictionary problem

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### Multiple pattern matching problem

- Given a set of pattern  $P = \{P_1, P_2, \dots, P_k\}$  and a text (databases)  $T$ , find all the occurrences of all the patterns.
- The sum of the lengths of patterns:  $m$ , length of text:  $n$ . Previous algorithms imply a search algorithm of  $\theta(m+kn)$  time.
- There are algorithms running in  $\theta(m+n+l)$  time, where  $l$  is the total number of occurrences of all the patterns
- Using keyword tree of  $P$
- The same idea as in KMP

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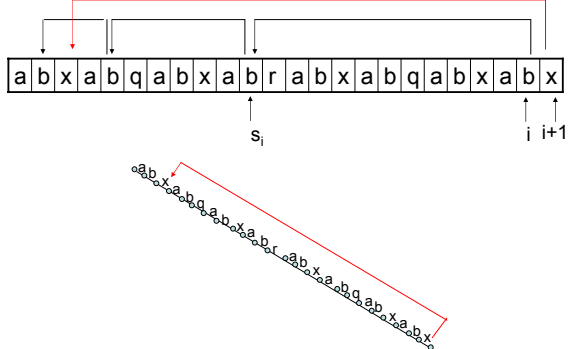
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### Failure function




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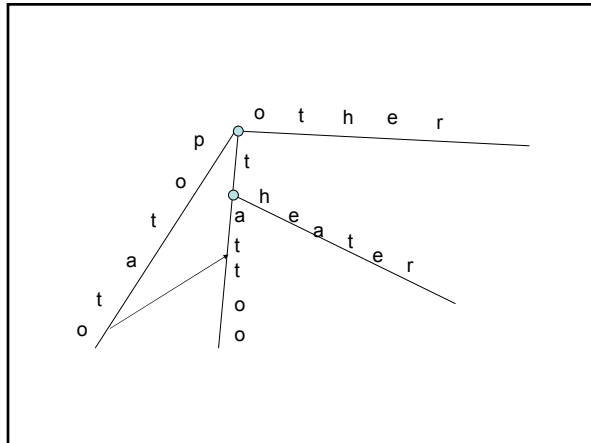
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- $L(v)$  denote the string from the root to  $v$  and  $lp(v)$  denote the length of the longest proper suffix of string  $L(v)$  that is a prefix of some pattern in  $P$ .
- $lp(v)$  for all the node  $v$  in  $K$  can be computed in linear time  $\theta(m)$
- Use the same failure links  $v \rightarrow n(v)$

[illegible][illegible]

- Create the keyword tree
- Compute the  $lp(v)$  and  $n(v)$  for each node  $v$  in the keyword tree
- Search the text against the tree, when a mismatch,  $T$  shifts  $lp(v)$  spaces and starts to compare from  $n(v)$
- Total running time  $\theta(m+n+I)$

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### Exact string matching applications

- Sequence-tagged-sites
- Exact string matching with wild cards
- Two-dimensional exact matching
- Regular expression pattern matching

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### Suffix tree

- Introduction
- Construction
- Applications
  
- Reading: Gusfield Ch5-7

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### Suffix tree

- Given a finite alphabet set  $\Sigma$ , a string S of length m, e.g.,  $S=abxabc$
- Suffix tree of S:
  - Rooted, directed
  - Edges labeled by **substrings** of S
  - Edges coming out of a node start with distinct letters
  - Exactly m leaves
  - **Leaf i spells out suffix  $S[i..m]$**

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## Keyword tree

- Keyword tree: given a set of pattern  $P = \{P_1, P_2, \dots, P_k\}$ , the keyword tree  $K$  is a tree:
  - Rooted, directed
  - Each edge is labeled with one letter
  - Edges coming out of a node have distinct labels
  - Every pattern is spelled out (map to one node)
  - Every leaf maps to some pattern

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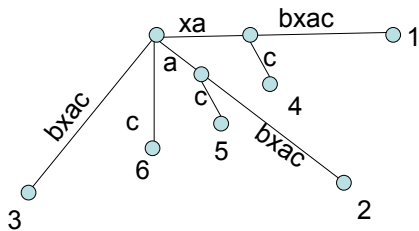
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## Suffix tree

Example: xabxac



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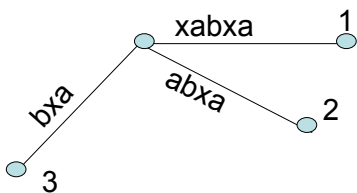
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## Suffix tree

- What if a suffix is a prefix of some other suffix? E.g. xabxa
- Solution: append a new letter \$
- Implicit suffix tree: if we don't require each leaf maps to one suffix.



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## First construction algorithm

- For a string S:
  - Assume no suffix is a prefix of some other suffix.
  - Make every suffix as a pattern  $P_i = S[i..m]$
  - Apply the linear time keyword tree construction algorithm
  - Concatenate “paths” into “edges”
- Running time:
  - Linear in the sum of the lengths of patterns  $\sum_i |P_i| = m(m+1)/2$
  - $\theta(m^2)$
  - Goal: design a linear time algorithm  $\theta(m)$

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## Why suffix tree

- 1<sup>st</sup> application: exact matching
- Suppose in  $\theta(n)$  time we can build the suffix tree for the text T
- Given any pattern P, at any time
  - Match letters of P along the suffix tree, until
  - Either no more matches are possible, P doesn't occur anywhere in T
  - Or P is exhausted: the labels of the leaves in side the subtree under the last matching edge are the starting positions of the occurrences.

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## 1<sup>st</sup> application

- Conclusion:
  - Exact string matching done in  $\theta(m+n+1)$  time
  - Exact multiple string matching done in  $\theta(m+n+1)$  time,
  - L the number of occurrences
- Other applications:
  - Multiple keyword search
  - Longest repeating substring
  - Longest common substring of two/more strings

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## Ukkonen's linear time construction

- Using implicit suffix tree of  $S[1..i]$ :  $T_i$ .
- Construct  $T_i$  incrementally:
  - From  $T_i$  to  $T_{i+1}$
  - Need to add  $S[i+1]$  to every suffix of  $S[1..i]$  and the empty string, there are  $i+1$  of them...
  - Append  $S[i+1]$  to suffix  $S[j..i]$  – becoming a suffix of  $S[j..(i+1)]$ ,  $j=1, 2, \dots, i, i+1$ .
  - Three suffix extension rules:

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## Suffix extension rules

- Let  $\beta$  denote the path of  $S[j..i]$
- 1.  $\beta$  ends at a leaf, append  $S[i+1]$  to the corresponding edge label
- 2. Some paths start from the end of  $\beta$ , but none of them starts with  $S[i+1]$ , add a new leaf edge, labeled by  $S[i+1]$
- 3. Some path from the end of  $\beta$  starts with  $S[i+1]$ , already in the tree, do nothing
- Straightforward implementation  $\theta(m^3)$

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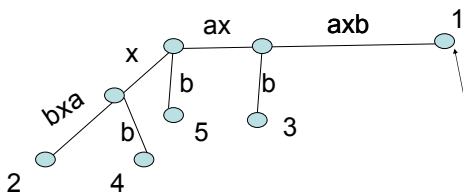
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## Ukkonen's algorithm, speedup

- How?
- Key thing to do: to locate the ending position of  $S[j..i]$  in  $T_i$
- Example:  $S[1..5] = axaxb$ , try adding  $S[6]=a$ .




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## Suffix links

- $j=1$ : easy (use a pointer pointing to the longest path in  $T_j$ ) append  $a$  to the edge
- Denote the leaf edge as  $(v, 1)$ 
  1. If  $v$  is the root,  $j=2$  is done straightforwardly
  2.  $v$  is not the root: there is another node, denoted as  $s(v)$ , such that if root to  $v$  spells out  $S[1..i]$ , then root to  $s(v)$  spells out  $S[2..i]$
  3. When we have the information on  $s(v)$ , continue search from it, not necessarily from the root again.
- Repeat for every  $j$

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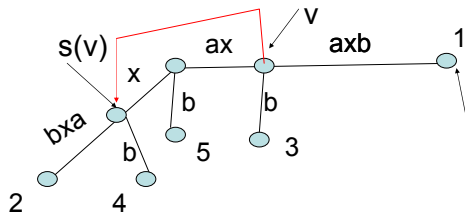
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## Suffix links

- Notes:
  - need to record  $s(v)$  if a new node is created
  - It doesn't give a "faster" algorithm




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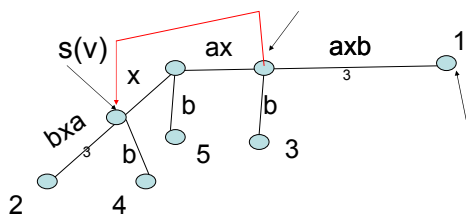
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## Trick1: skip/count

- Key: record the length of each edge,
- When search the path, by using the length information, for each step, we can go node by node, not character by character; only constant number of comparison at each node (assume constant alphabet set)




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## Analysis of trick 1

- Every node has a depth, i.e., the number of nodes on the path from the root
- The depth of  $v$  is at most one greater than the depth of  $s(v)$ ,
- During the entire phase constructing  $T_{i+1}$  from  $T_i$  ( $1 \leq j \leq i+1$ ): decrease node depth at most  $2m$  (each  $j$ , decrease 1 to find  $v$ , and decrease 1 to find  $s(v)$ ;  $i \leq m$ )
- The total number of node length that could be increased during a phase is bounded by the number of decrement ( $2m$ ) and the maximum length of a path ( $m$ )
- So construction done in  $O(m)$
- Applying suffix link and trick 1 gives an  $O(m^2)$  time suffix tree building algorithm

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## Two observations

- Observation1: The space for the total number of letters in the edge labels could reach  $\theta(m^2)$
- An alternate way to represent labels is necessary: Use position intervals  $[start, end]$  to represent label  $S[start..end]$
- Observation2: When  $S[j..i]$  doesn't end at a leaf and there is an extending edge whose label starts with  $S[i+1]$ , we are done for  $j$ , we are done for the phase  $i$  ( $S[j+1, i+1]$ ,  $S[j+2, i+1]$ , ...,  $S[i, i+1]$ ,
- Trick 2: whenever this happens,  $T_{i+1}$  is built

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## Last trick

- Observation 3: once a leaf, always a leaf. If a leaf is created and labeled by an index  $j$ , rule 1 will always apply.
- Trick 3: use a global parameter  $e$  to denote the last position thus to skip the extensions (only need to update  $e$  once per iteration).

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### Recall: Ukkonen's linear time construction

- Using implicit suffix tree of  $S[1..i]$ :  $T_i$ .
- Construct  $T_i$  incrementally:
  - From  $T_i$  to  $T_{i+1}$
  - Need to add  $S[i+1]$  to every suffix of  $S[1..i]$  and the empty string, there are  $i+1$  of them...
  - Append  $S[i+1]$  to suffix  $S[j..i]$  – becoming a suffix of  $S[1..(i+1)]$ ,  $j=1, 2, \dots, i, i+1$ .
  - Three suffix extension rules:

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### Recall: Suffix extension rules

- Let  $\beta$  denote the path of  $S[j..i]$
- 1.  $\beta$  ends at a leaf, append  $S[i+1]$  to the corresponding edge label
- 2. Some paths start from the end of  $\beta$ , but none of them starts with  $S[i+1]$ , add a new leaf edge, labeled by  $S[i+1]$
- 3. Some path from the end of  $\beta$  starts with  $S[i+1]$ , already in the tree, do nothing

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### Time complexity: amortized analysis

- Increment  $e$  to skip the first  $j^*$  (from last phase) extensions. (Skip some  $j$  at the beginning)
- Apply trick 1 to continue until trick 2 can be applied at  $j^{\text{th}}$  extension, (skip some  $j$  at last)
  - Set  $j^*=j-1$ : update  $j^*$  for the next phase use
- Next phase we can skip the first  $j^*$  extensions...
- Every two consecutive phases overlap at most 1 index
- Linear time algorithm!
- ( $j^*$  of  $(i+1)^{\text{th}}$  run is the previous position that rule 3/trick 2 applied in  $i^{\text{th}}$  run.

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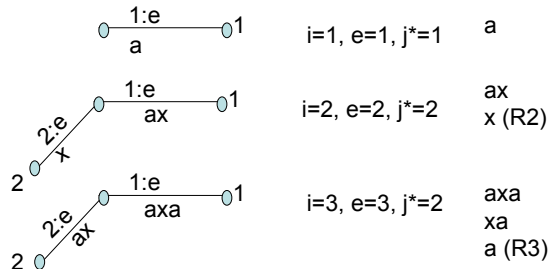
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### Example

- $S[1..6]=\text{axaxba}$




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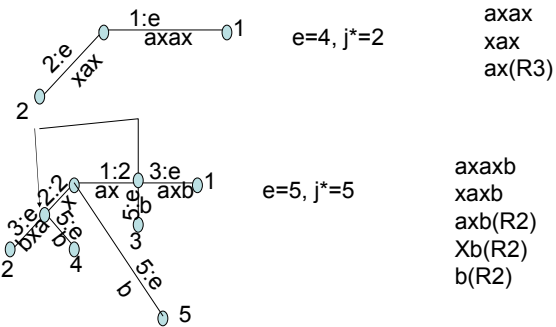
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### Example

- $S[1..6]=\text{axaxba}$




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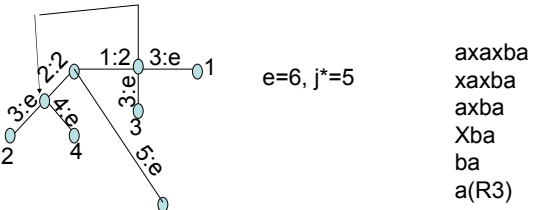
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### Example

- $S[1..6]=\text{axaxba}$




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### The true suffix tree

- Append \$ to S and execute on  $T_m$
- Correctness: every suffix is spelled out by some root-to-leaf path, no suffix is a prefix of some other suffix
- Generalized suffix tree for a set of strings:
  - Concatenate strings into one, by adding some extra letters, but some synthetic suffixes
  - Build suffix tree for one string, then on top of it build for another, then on top of it, build for another...

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### Applications

- Exact pattern search
- Longest common substring
- Tandem repeat
- Suffix array
- ...

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