## Information Retrieval and Organisation



## Suffix Trees

adapted from
http://www.math.tau.ac.il/~haimk/seminar02/suffixtrees.ppt

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## Trie

- A tree representing a set of strings



## Trie

- Assume no string is a prefix of another

1) Each edge is labeled by a letter.
2) No two edges outgoing from the same node are labeled the same.
3) Each string corresponds to a leaf.


## Compressed Trie

- Compress unary nodes, label edges by strings



## Suffix Tree

- Given a string $s$, a suffix tree of $s$ is a compressed trie of all suffixes of $s$.
- To make these suffixes prefix-free we add a special character, say $\$$, at the end of $s$.


## Suffix Tree

- For example, let $s=$ abab, a suffix tree of $s$ is a compressed trie of all suffixes of abab\$.


Note that a suffix tree has $\mathrm{O}(n)$ nodes $n=|\mathbf{s}|$. Why?

## Suffix Tree Construction

The trivial algorithm

Put the largest suffix in



Put the suffix bab\$ in



Put the suffix ab\$ in



Put the suffix b\$ in



Put the suffix \$ in



We will also label each leaf with the starting point of the corresponding suffix


## Suffix Tree Construction

- The trivial algorithm takes $\mathrm{O}\left(n^{2}\right)$ time.
- It is possible to build a suffix tree in $\mathrm{O}(n)$ time using Ukkonen's algorithm.
- But, how come? Does it take O(n) space?
- To use only O(n) space, encode the edge-labels as (beginning-position, end-position).


## Consider the string aaaaaabbbbbb\$



## Consider the string aaaaaabbbbbb\$



## Consider the string aaaaaabbbbbb\$



## Suffix Tree Applications

- What Can We Do with It?
- Exact String Matching
- Exact Set Matching
- The Substring Problem for a Database of Patterns
- Longest Common Substring of Two Strings
- Recognising DNA Contamination
- Common Substring of More Than Two Strings


## Exact String Matching

- Given text $T(|T|=n)$, pre-process it such that when a pattern $P(|P|=m)$ arrives you can quickly decide when it occurs in $T$.
- We may also want to find all occurrences of $P$ in T.


## Exact String Matching

- In pre-processing, we just build a suffix tree in $\mathrm{O}(n)$ time



## Exact String Matching

- Given a pattern $P=\mathrm{ab}$ we traverse the tree according to the pattern.
- If we do not get stuck traversing the pattern then the pattern occurs in the text, otherwise it does not.
- Each leaf in the subtree below the node we reach corresponds to an occurrence.
- By traversing this subtree we get all $k$ occurrences in $\mathrm{O}(n+k)$ time.


## Exact String Matching

- How to match a pattern (query) against a database of strings (documents)?


## Generalized Suffix Tree

- Given a set of strings S, the generalized suffix tree of $S$ is a compressed trie of all suffixes of each $s \in S$.
- To make these suffixes prefix-free we add a special char, say $\$$, at the end of $s$.
- To associate each suffix with a unique string in $S$, add a different special char to each s.
- Each leaf node needs to be labelled by the document id together with the suffix position.


## Generalized Suffix Tree

- For example, Let $s_{1}=a b a b$ and $s_{2}=a a b$, here is a generalized suffix tree for $s_{1}$ and $s_{2}$.



## Longest Common Substring

- Given two strings $s_{1}$ and $s_{2}$, we build their generalized suffix tree.
- Every node with a leaf descendant from string $s_{1}$ and a leaf descendant from string $s_{2}$ represents a maximal common substring and vice versa.
- Find such node with largest "string depth".


## Lowest Common Ancestor

- A lot more can be gained from the suffix tree, if we pre-process it so that we can answer LCA queries on it in constant time.



## Lowest Common Ancestor

- Why? The LCA of two leaves represents the longest common prefix (LCP) of these 2 suffixes



## Finding Maximal Palindromes

- A palindrome: cbaabc, caabaac, ...
- To find all palindromes in a string $s$ (of length $m$ ), we build a generalized suffix tree for the string $s$ and the reversed string $s^{r}$.
- The palindrome with centre between $i-1$ and $i$ is the LCP of the suffix at position $i$ of $s$ and the suffix at position $m-i$ of $s^{r}$.


## Finding Maximal Palindromes

- For example, consider the string cbaaba.
- Prepare a generalized suffix tree for $s=$ cbaaba\$ and $s^{r}=$ abaabc\#
- For every $i$ find the LCA of the suffix $i$ of $s$ and the suffix $m-i$ of $s^{r}$.
- All palindromes can be identified in linear time.

Let $s=$ cbaaba $\$$ then $s^{r}=$ abaabc\#


## Suffix Tree Drawbacks

- It is $\mathrm{O}(n)$ but the constant is quite big.
- It consume a lot of space.
- Notice that if we indeed want to traverse an edge in $\mathrm{O}(1)$ time then we need an array (of pointers) of size $|\Sigma|$ in each node, where $\Sigma$ is the alphabet.


## Suffix Array

- It is much simpler and easier to implement.
- Compared with suffix trees, we lose some functionality, but we save space.


## Suffix Array

- For example, let s = abab
- Sort the suffixes lexicographically: ab, abab, b, bab
- The suffix array gives the indices of the suffixes in sorted order



## Suffix Array Construction

- The trivial algorithm
- Quicksort
- The linear time algorithm
- Build a suffix tree in $\mathrm{O}(n)$ time first, and then traverse the tree in in-order, lexicographically picking edges outgoing from each node, and fill the suffix array.
- It can also be built in $\mathrm{O}(n)$ time directly.


## Exact String Matching

- How do we search for a pattern $P$ in the text $T$, using the suffix array of $T$ ?
- If $P$ occurs in $T$, then all its occurrences are consecutive in the suffix array.
- So we can do two binary searches on the suffix array: the first search locates the starting position of the interval, and the second one determines the end position.
- It takes $\mathrm{O}(m \log (n))$ time, as a single suffix comparison needs to compare up to $m$ characters.


## Exact String Matching

- It is also possible to do it in $\mathrm{O}(m+\log (n))$ with an additional array of LCP.
- Manber \& Myers (1990)
$T=$ mississippi
$P=$ issa

| $L \longrightarrow$ | 10 | i |
| :---: | :---: | :---: |
|  | 7 | ippi |
|  | 4 | issippi |
|  | 1 | ississippi |
|  | 0 | mississippi |
| $\mathbf{M} \longrightarrow$ | 9 | pi |
|  | 8 | ppi |
|  | 6 | sippi |
|  | 3 | sisippi |
|  | 5 | ssippi |
| $\mathrm{R} \longrightarrow$ | 2 | ssissippi |

