

Part I

Suffix trees and their uses

Introduction to suffix trees

A suffix tree is a data structure that exposes the internal structure of a string in a deeper way than does the fundamental preprocessing discussed in Section ???. Suffix trees can be used to solve the exact matching problem in linear time (achieving the same worst case bound that the Knuth-Morris-Pratt and the Boyer-Moore algorithms achieve), but their real virtue comes from their use in linear time solutions to many string problems more complex than exact matching. Moreover (as we will detail in Chapter ???), suffix trees provide a *bridge* between *exact* matching problems, the focus of part I, and *inexact* matching problems that are the focus of Part III.

The classic application for suffix trees is the *substring problem*. One is first given a text T of length m . After $O(m)$, or linear, preprocessing time, one must be prepared to take in *any unknown string* S of length n and in $O(n)$ time either find an occurrence of S in T or determine that S is not contained in T . That is, the allowed preprocessing takes time proportional to the length of the text, but thereafter, the search for S must be done in time proportional to the length of S , *independent* of the length of T . These bounds are achieved with the use of a suffix tree. The suffix tree for the text is built in $O(m)$ time during a preprocessing stage; thereafter whenever a string of length $O(n)$ is input, the algorithm searches for it in $O(n)$ time using that suffix tree.

The $O(m)$ preprocessing and $O(n)$ search result for the substring problem is very surprising and extremely useful. In typical applications, a long sequence of requested strings will be input after the suffix tree is built, so the linear time bound for each search is important. That bound is *not* achievable by the Knuth-Morris-Pratt or Boyer-Moore methods – those methods would preprocess each requested string on input, and then take $\Theta(m)$ (worst case) time to search for the string in the text. Since m may be huge compared to n , those algorithms would be impractical on any but trivial sized texts.

Often the text is a fixed *set* of strings, for example a collection of STSs or ESTs (see Sections ?? and ??), so that the substring problem is to determine whether the input string is a substring of any of the fixed strings. Suffix trees work nicely to efficiently solve this problem as well. Superficially, this case of multiple text strings resembles the *dictionary* problem discussed in the context of the Aho-Corasick algorithm. So it is natural to expect that the Aho-Corasick algorithm could be applied. However, the Aho-Corasick method does not solve the substring problem in the desired time bounds, because it will only determine if the new string is a *full* string in the dictionary, not whether it is a substring of a string in the dictionary.

After presenting the algorithms, several applications and extensions will be discussed in Chapter ???. Then a remarkable result, *the constant time least common ancestor method*, will be presented in Chapter ???. That method greatly amplifies

the utility of suffix trees, as will be illustrated by additional applications in Chapter ???. Some of those applications provide a bridge to inexact matching, and more applications of suffix trees will be discussed in Part III where the focus is on inexact matching.

1 A short history

The first linear time algorithm for constructing suffix trees was given by Weiner [?] in 1973, although he called his tree a position tree. A different, more space efficient algorithm to build suffix trees in linear time was given by McCreight [?] a few years later. Recently, Ukkonen [?] developed a conceptually different linear time algorithm for building suffix trees which has all the advantages of McCreight's algorithm (and when properly viewed can be seen as a variant of McCreight's algorithm) but allows a much simpler explanation.

Although more than twenty years have passed since Weiner's original result (which Knuth is claimed to have called "the algorithm of 1973" [?]), suffix trees have not made it into the mainstream of computer science education, and have generally received less attention and use than might have been expected. This is probably because the two original papers of the 1970's have a reputation for being extremely difficult to understand. That reputation is well deserved but unfortunate, because the algorithms, although non-trivial, are not more complicated than many widely taught methods. And, when implemented well, the algorithms are practical and allow efficient solutions to many complex string problems. We know of no other single data structure (other than those essentially equivalent to suffix trees) that allows efficient solutions to such a wide range of complex string problems.

Chapter ??? fully develops the linear time algorithms of Weiner and Ukkonen, and then briefly mentions the high level organization of McCreight's algorithm and its relationship to Ukkonen's algorithm. Our approach is to introduce each algorithm at a high level, giving simple, *inefficient* implementations. Those implementations are then incrementally improved to achieve linear running times. We believe that the expositions and analyses given here, particularly for Weiner's algorithm, are much simpler and clearer than in the original papers, and hope that these expositions result in a wider use of suffix trees in practice.

Note added March 2006: Today the simplest way to show that suffix trees can be built in linear time, and an approach that is efficient in practice, is to show how to build a suffix array and auxiliary longest common prefix array, in linear time, and then how to build the full suffix tree from those arrays in linear time.

To avoid this problem, we assume (as was true in Figure 1) that the last character of S appears nowhere else in S . Then, no suffix of the resulting string can be a prefix of any other suffix. To achieve this in practice, we can add a character to the end of S that is not in the alphabet that string S is taken from. In this book we use $\$$ for the “termination” character. When it is important to emphasize the fact that this termination character has been added, we will write it explicitly as in $S\$$. But much of the time, this reminder will not be necessary and, unless explicitly stated otherwise, every string S is assumed to be extended with the termination symbol $\$$, even if the symbol is not explicitly shown.

A suffix tree is related to the keyword tree (without backpointers) considered in Section ???. Given string S , if set \mathcal{P} is defined to be the m suffixes of S , then the suffix tree for S can be obtained from the keyword tree for \mathcal{P} by merging any path of non-branching nodes into a single edge. The simple algorithm given in Section ??? for building keyword trees could be used to construct a suffix tree for S in $O(m^2)$ time, rather than the $O(m)$ bound we will establish.

Definition The *label of a path* from the root that ends at a *node* is the concatenation, in order, of the substrings labeling the edges of that path. The *path-label of a node* is the label of the path from the root of \mathcal{T} to that node.

Definition For any node v in a suffix tree, the *string-depth* of v is the number of characters in v 's label.

Definition A path that ends in the middle of an edge (u, v) splits the label on (u, v) at a designated point. Define the label of such a path as the label of u concatenated with the characters on edge (u, v) down to the designated split point.

For example, in Figure 1 string xa labels the internal node w (so node w has path-label xa), string a labels node u , and string $xabx$ labels a path that ends inside edge $(w, 1)$, i.e. inside the leaf edge touching leaf 1.

3 A motivating example

Before diving into the details of the methods to construct suffix trees, let's look at how a suffix tree for a string is used to solve the *exact match* problem: given a pattern P of length n and a text T of length m , find all occurrences of P in T in $O(n + m)$ time. We have already seen several solutions to this problem. Suffix trees provide another approach.

Build a suffix tree \mathcal{T} for text T in $O(m)$ time, Then match the characters of P along the unique path in \mathcal{T} until either P is exhausted or no more matches are possible. In the latter case P does not appear anywhere in T . In the former case, every leaf in the subtree below the point of the last match is numbered with a starting location of P in T , and every starting location of P in T numbers such a leaf.

The key to understanding the former case (when all of P matches a path in T) is to note that P occurs in T starting at position j if and only if P occurs as a prefix of $T[j..m]$. But that happens if and only if string P labels an initial part of the path from the root to leaf j . It is that initial path that will be followed by the matching algorithm.

The matching path is unique because no two edges out of a common node can have edge-labels beginning with the same character. And, because we have assumed a finite alphabet, the work at each node is constant and so the time to match P to a path is proportional to the length of P .

For example, Figure 2 shows a fragment of the suffix tree for string $T = awyawxawxz$. Pattern $P = aw$ appears three times in T starting at locations 1, 4 and 7. Pattern P matches a path down to the point shown by an arrow, and as required, the leaves below that point are numbered 1, 4 and 7.

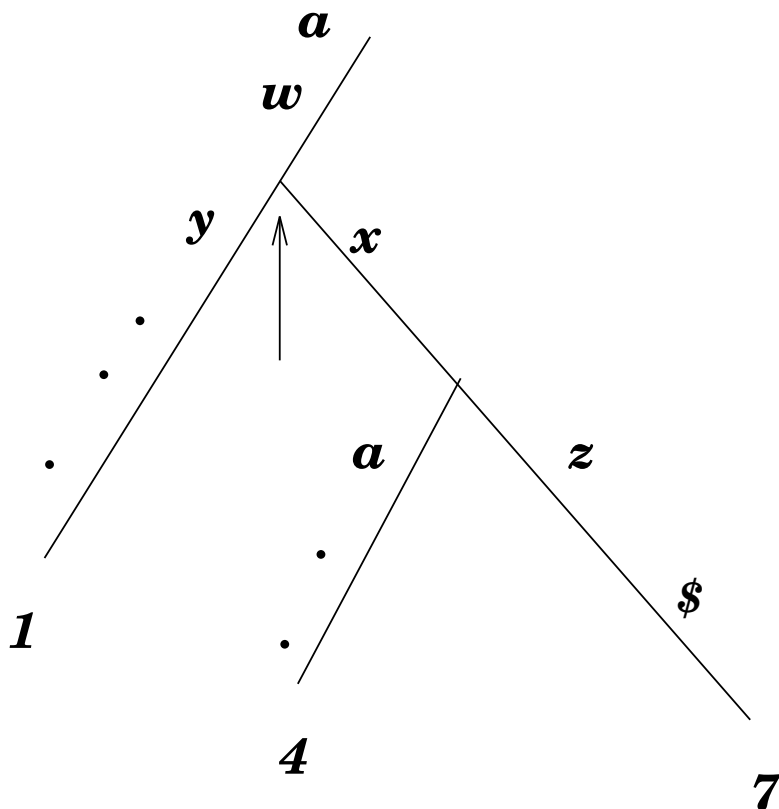


Figure 2: Three occurrences of aw in $awyawxawxz$. Their starting positions number the leaves in the subtree of the node with path-label aw .

If P fully matches some path in the tree, the algorithm can find all the starting positions of P in T by traversing the subtree below the end of the matching path, collecting position numbers written at the leaves. So all occurrences of P in T can be found in $O(n + m)$ time. This is the same overall time bound achieved by several

algorithms considered in Part I, but the distribution of work is different. Those earlier algorithms spend $O(n)$ time for preprocessing P , and then $O(m)$ time for the search. In contrast, the suffix tree approach spends $O(m)$ preprocessing time, and then $O(n + k)$ search time, where k is the number of occurrences of P in T .

To collect the k starting positions of P , traverse the subtree at the end of the matching path using any linear time traversal (depth-first say), and note the leaf numbers encountered. Since every internal node has at least two children, the number of leaves encountered is proportional to the number of edges traversed, so the time for the traversal is $O(k)$, even though the the total string-depth of those $O(k)$ edges may be arbitrarily larger than k .

If only a single occurrence of P is required, and the preprocessing is extended a bit, then the search time can be reduced from $O(n + k)$ to $O(n)$ time. The idea is to write at each node one number (say the smallest) of a leaf in its subtree. This can be achieved in $O(m)$ time in the preprocessing stage by a depth-first traversal of \mathcal{T} . The details are straightforward and are left to the reader. Then, in the search stage, the number written on the node at or below the end of the match gives one starting position of P in T .

In Section ?? we will again consider the relative advantages of methods that preprocess the text versus methods that preprocess the pattern(s). Later, in Section ?? we will also show how to use a suffix tree to solve the exact matching problem using $O(n)$ preprocessing and $O(m)$ search time, achieving the same bounds as in the algorithms presented in Part I.

4 A naive algorithm to build a suffix tree

To further solidify the definition of a suffix tree and develop the reader's intuition, we present a straightforward algorithm to build a suffix tree for string S . This naive method first enters a single edge for suffix $S[1..m]$ (the entire string) into the tree, then it successively enters suffix $S[i..m]$ into the growing tree, for i increasing from 2 to m . We let N_i denote the intermediate tree that encodes all the suffixes from 1 to i .

In detail, tree N_1 consists of a single edge between the root of the tree and a leaf labeled 1. The edge is labeled with the string S . Tree N_{i+1} is constructed from N_i as follows: Starting at the root of N_i find the longest path from the root whose label matches a prefix of $S[i+1..m]$. This path is found by successively comparing and matching characters in suffix $S[i+1..m]$ to characters along a unique path from the root, until no further matches are possible. The matching path is unique because no two edges out of a node can have labels which begin with the same character. At some point, no further matches are possible because no suffix of S is a prefix of any other suffix of S . When that point is reached, the algorithm is either at a node, w say, or it is in the middle of an edge, (u, v) say, then it breaks edge (u, v) into two edges by inserting a new node, called w , just after the

last character on the edge that matched a character in $S[i + 1..m]$, and just before the first character on the edge that mismatched. The new edge (u, w) is labeled with the part of the (u, v) label that matched with $S[i + 1..m]$, and the new edge (w, v) is labeled with the remaining part of the (u, v) label. Then (whether a new node w was created, or already existed at the point where the match ended), the algorithm creates a new edge $(w, i + 1)$ running from w to a new leaf labeled $i + 1$, and it labels the new edge with the unmatched part of suffix $S[i + 1..m]$.

The tree now contains a unique path from the root to leaf $i + 1$, and this path is labeled with the string $S[1 + 1..m]$. Note that all edges out of the new node w have labels which begin with different first characters, and so it follows inductively that no two edges out of a node have labels with the same first character.

Assuming, as usual, a bounded-size alphabet, the above naive method takes $O(m^2)$ time to build a suffix tree for the string S of length m .