BMI/CS 776 Lecture #17: Pattern Matching Suffix trees

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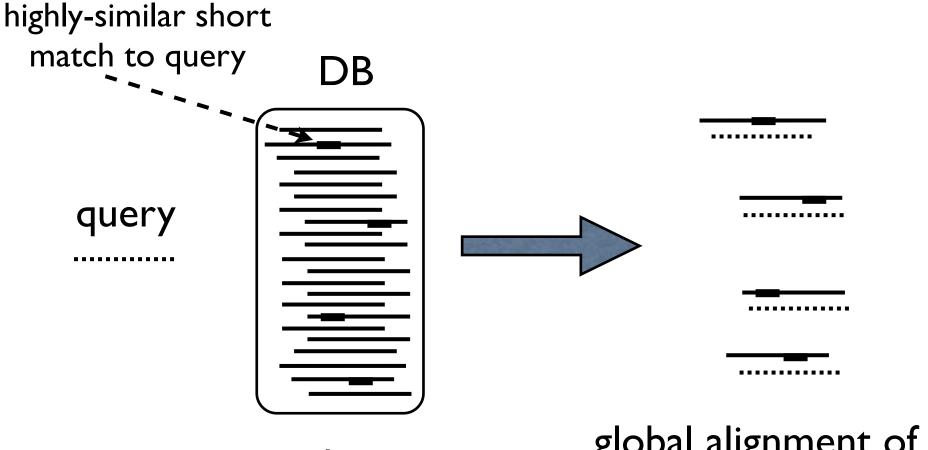
Alignment vs pattern matching

- Global sequence alignment
 - Input: n ≥ 2 relatively short sequences
 - Homology assumptions: homologous along entire length, colinear
 - Goal: determine homologous positions
- Pattern matching
 - Input: n ≥ I sequences (short or long)
 - Homology assumptions: none
 - Goal: find short exact/inexact substring (local) matches between or within input sequences

Why pattern matching?

- Applications:
 - Database search short query, large DB
 - Alignment of long sequences exact global alignment no longer feasible
 - Alignment of rearranged/duplicated sequences no longer global alignment
- Key idea:
 - Short local matches can seed longer global alignments

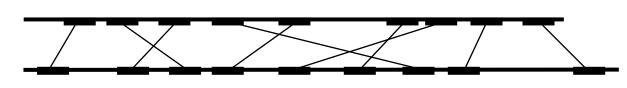
Database search



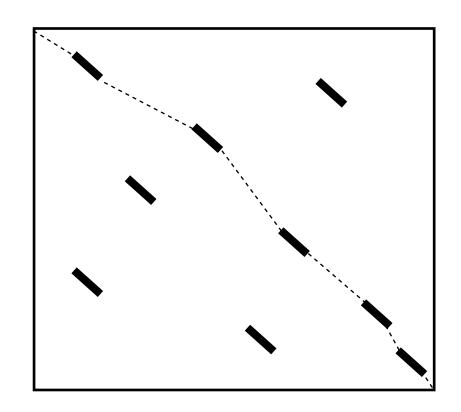
pattern matching

global alignment of each candidate to query

Global alignment of long sequences

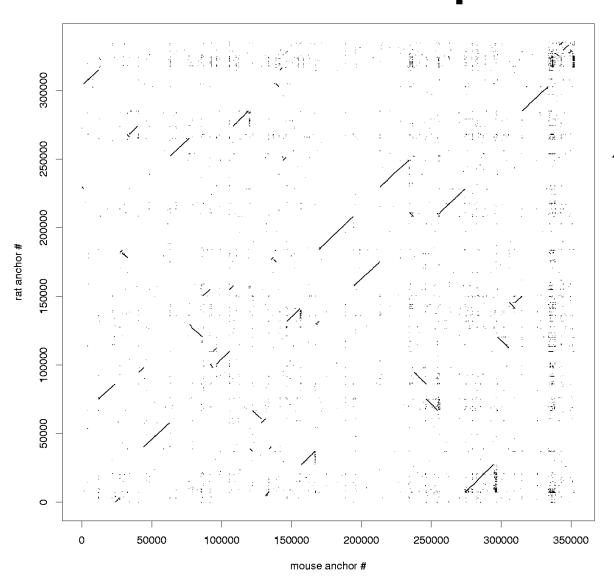


Find seed matches using pattern matching



Sparse dynamic programming with seed matches

Alignment of rearranged sequences



whole-genome

dot plot

Each dot represents a seed match between genome sequences

Suffix trees

- Very important for pattern matching
- Substring problem:
 - Given text T of length m
 - Preprocess T in O(m) time
 - Such that, given query string S of length n,
 find occurrence (if any) of S in T in O(n) time
- Suffix trees solve this problem, and many more

History of suffix trees

- Weiner (1973) First linear time algorithm for suffix tree construction
 - "the algorithm of 1973" (Knuth)
- McCreight (1976) Space efficient version of algorithm

this lecture

 Ukkonen (1995) - Simple, elegant algorithm, with "online" property

Suffix tree definition

- A suffix tree T for a string S of length m is tree with the following properties:
 - rooted and directed
 - m leaves, labeled 1 to m

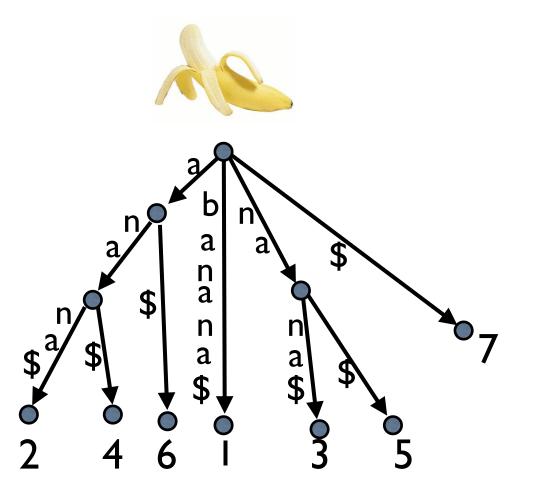
key property



- Each edge labeled by a substring of S
- Concatenation of edge labels on path from root to leaf i is suffix i of S (we will denote this by $S_{i...m}$)
- Each internal non-root node has at least two children
- Edges out of a node must begin with different characters

Suffix tree example

- *S* = "banana\$"
- Add '\$' to end so that suffix tree exists

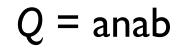


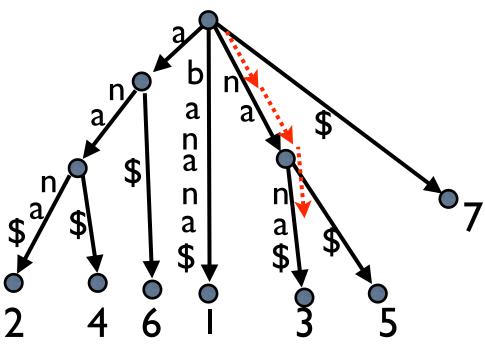
Solving the substring problem

- Assume we have suffix tree T
- FindMatch(*Q*,*T*):
 - Follow (unique) path down from root of T according to characters in Q
 - If all of Q is found to be a prefix of such a path
 - return label of some leaf below this path
 - else, return no match found

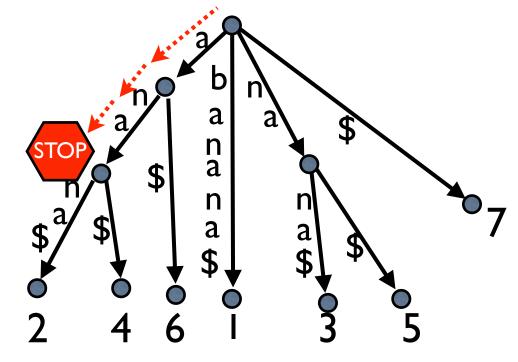
Solving the substring problem







return 3



return no match found

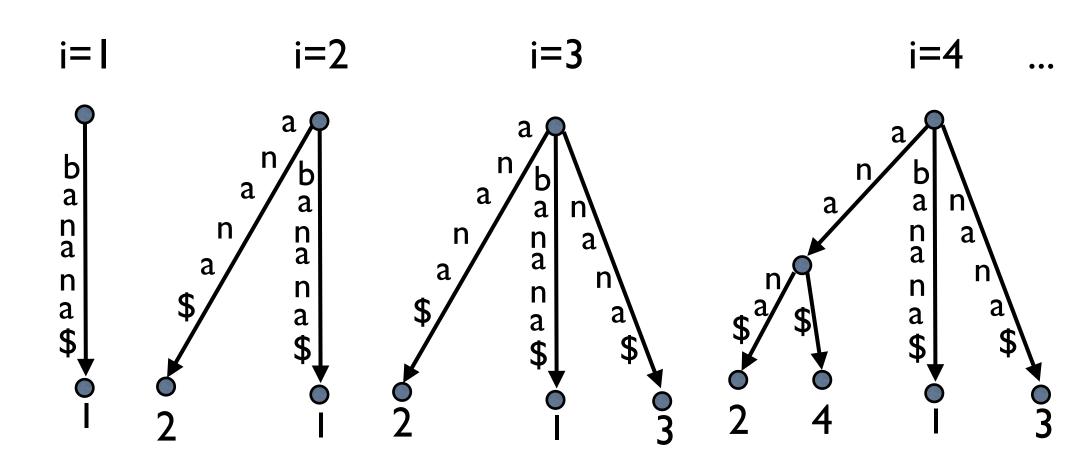
Runtime of substring problem with suffix tree

- Finite alphabet: O(1) work at each node
- Edges out of each node start with unique characters: unique path from root
- Size of tree below end of path: O(k), k = number of suffixes starting with Q
- O(n + k) time to report all k matching substrings
- O(n) to report just one with an additional trick

Naive suffix tree building

- Now we need a O(m) time algorithm for building suffix trees
- Naive algorithm is $O(m^2)$:
 - $T \leftarrow$ empty tree
 - For i from 1 to m:
 - Add suffix $S_{i...m}$ to T by finding longest matching prefix of $S_{i...m}$ in T and branching from there

O(m²) suffix tree building

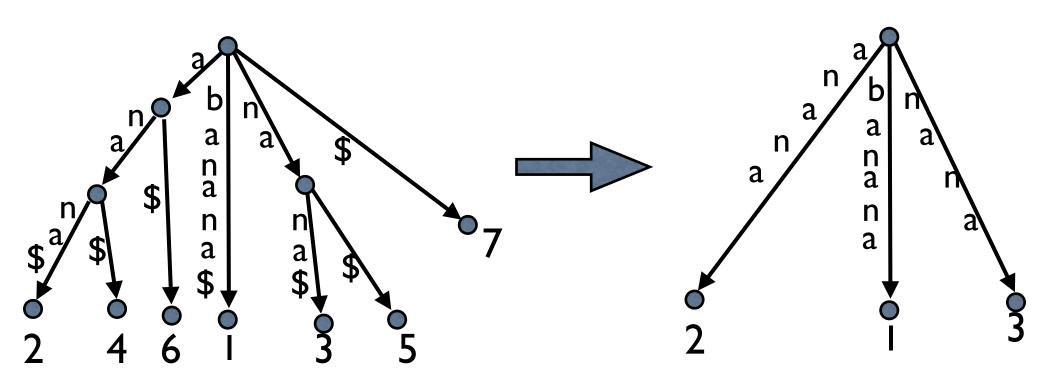


Ukkonen's O(m) algorithm

- On-line algorithm
 - Builds implicit suffix tree for each prefix of string S
 - Implicit suffix tree of S_{1...i} denoted I_i
 - Builds I_1 , then I_2 from I_1 ,..., then I_m from I_{m-1}
- Basic algorithm is O(m³), but with a series of tricks, it is O(m)

Implicit suffix tree

- Suffix tree → implicit suffix tree
 - remove \$ characters from labels
 - remove edges with empty labels
 - remove internal nodes without two children

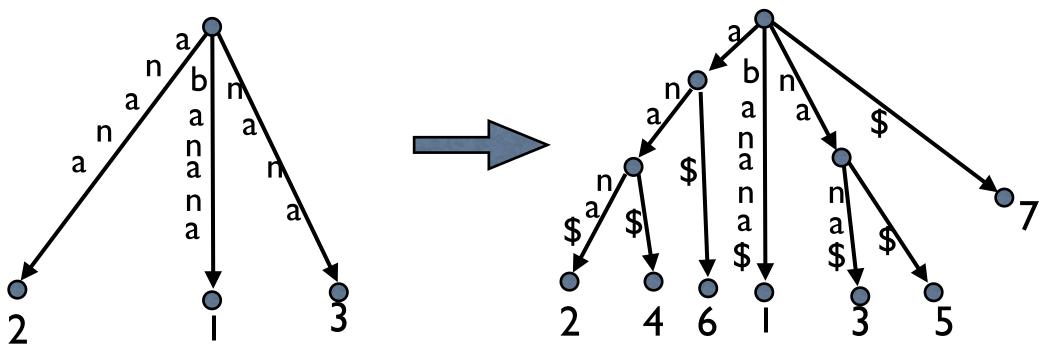


Ukkonen's algorithm overview

- Construct I₁
- For *i* from 1 to m 1:
 - For j from I to i + I:
 - Find end of path from root labeled $S_{j...i}$
 - Add character S_{i+1} to the end of this path in the tree, if necessary (suffix extension)

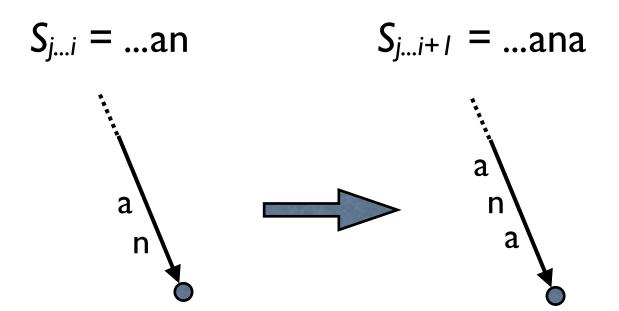
Conversion to suffix tree

- Convert implicit suffix tree at end of algorithm into true suffix tree
- Simply run algorithm for one more iteration with \$ final character
- Traverse tree to label leaf edges with positions



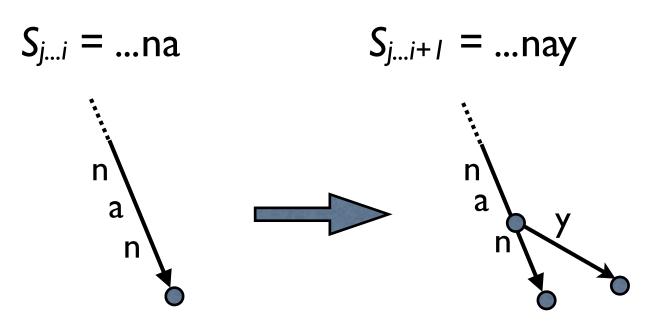
Suffix extension rule I

I. If path $S_{j...i}$ in tree ends at leaf, add character S_{i+1} to end of label of edge into leaf



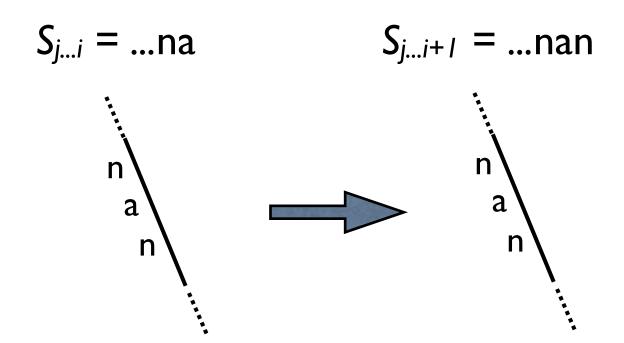
Suffix extension rule II

II. If there are paths continuing from path $S_{i...j}$ in the tree, but none starting with S_{i+1} , then create a new leaf edge with label S_{i+1} at the end of path $S_{i...j}$ (creating a new internal node if $S_{i...j}$ ends in the middle of an edge)



Suffix extension rule III

III. If there are paths continuing from path $S_{i...j}$ in the tree, and one starts with S_{i+1} , then do nothing

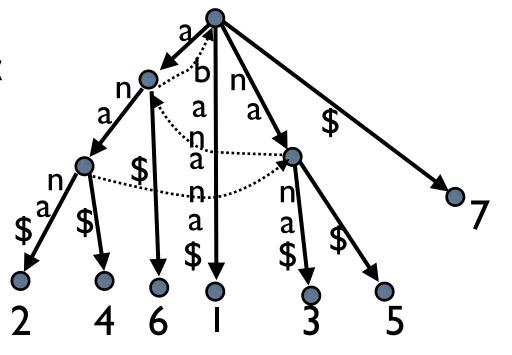


Suffix links

- How to find end of each suffix $S_{j...i}$?
- Could search down tree in O(i-j+1) time
 →O(m³) time total
- Better: create links between nodes corresponding to ends of similar suffixes
- With some additional tricks, get runtime to O(m²) time total

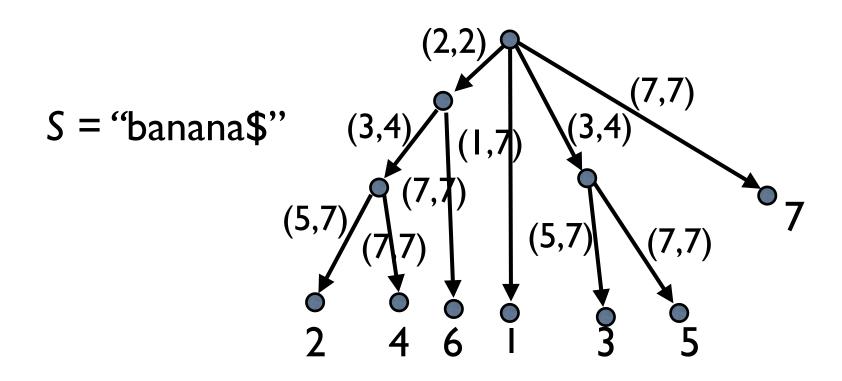
Suffix link definition

- A suffix link is a pointer from an internal node v to another node s(v) where
 - x is a character, α is a substring (possibly empty)
 - v has path-label $x\alpha$
 - s(v) has path-label α



Edge-label compression

- Label edges with pair of indices into string rather than with explicit substring
- Makes space requirement only O(m)



Final runtime

- With a few more tricks and implementation details, Ukkonen's algorithm runs in time O(m)
- More details found in (Ukkonen, 1995) or book by Dan Gusfield (Gusfield, 1997)