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Lecture 19: Text Indexing I

COMP526: Efficient Algorithms

Updated: December 5, 2024

Will Rosenbaum University of Liverpool

Announcements

- 1. Quiz 07 on Error Correcting Codes
 - Complete by 11:59pm, Friday 06 November
- 2. Grading is slow (sorry)
 - Programming assignment 1 grades next week
- 3. Last lectures:
 - Text indexing (Today and next Tuesday)
 - Final review (next Thursday)
- 4. Attendance Code:

Meeting Goals

- 1. Introduce and analyze Parallel MergeSort
- 2. Introduce the text indexing problem
- 3. Define the trie data structure
- 4. Define suffix trees
- 5. Describe applications of suffix trees

Parallel MergeSort

Last Time

Parallel Algorithms!

- PRAM model
 - Unlimited parallel processing elements (PEs)
- Brent's Theorem: span *T* and work *W* with unlimited PEs
 - \implies span O(T + W/p) and work O(W) with p PEs
- Parallel string matching with span T = O(m) and work W = O(n)
- Sorting networks
 - span $T = O(\log^2 n)$ and work $W = O(n\log^2 n)$
 - limited to specialized hardware and/or small arrays

Parallel Divide & Conquer?

Observation. The Divide & Conquer strategy can lend itself well to parallelism:

- 1. Divide problem into sub-tasks
- 2. Solve the subtasks
- 3. Merge solutions of the subtasks

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- 3. Merge solutions of the subtasks (...?)
 - How to parallelize this?

Parallel MergeSort?

Revisited: MERGESORT

1: **procedure** MERGESORT(A, i, k) if i < k then 2: $j \leftarrow \lfloor (i+k)/2 \rfloor$ 3: MERGESORT(A, i, j)4: MERGESORT(A, j + 1, k) 5: $B \leftarrow \text{COPY}(A, i, j)$ 6: $C \leftarrow \text{COPY}(A, j+1, k)$ 7: MERGE(B, C, A, i)8: end if 9: 10: end procedure

Parallel MergeSort?

PollEverywhere

What is the span of MergeSort with parallel recursive calls and sequential merges?



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- 1: **procedure** MERGESORT(*A*, *i*, *k*)
- 2: **if** *i* < *k* **then**
- 3: $j \leftarrow \lfloor (i+k)/2 \rfloor$
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- 5: MERGESORT(A, j+1, k)
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 - # in x's sub-array
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Idea.

• In x's own sub-array, just use x's index!

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- For the other sub-array, use binary search!

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Idea.

- In *x*'s own sub-array, just use *x*'s index!
- For the other sub-array, use binary search!
- **Parallelize:** do each *x* in parallel!

- 1: **procedure** PARALLELMERGE(*A*[*l..m*), *A*[*m..r*), *B*)
- 2: **for** $i = l, \dots, m-1$ **in parallel do**
- 3: $k \leftarrow (i l) + \text{BINARYSEARCH}(A[m..r), A[i])$
- 4: $B[k] \leftarrow A[i]$
- 5: **end for**
- 6: **for** j = m, m+1, ..., r-1 **in parallel do**
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 - Θ(log *n*)
- What is the **work** of PARALLELMERGE?
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Overall Procedure

- 1. Split (sub)array in half
- 2. Parallel recursive MergeSorts
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Improvements. Merge can be improved to $\Theta(n)$ work! (but it's complicated)

Concluding Thoughts

Parallelism is Necessary

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Parallelism is Subtle

- Reasoning about parallel programs is hard
- Writing correct parallel programs is hard
- Idealized models abstract away many challenges
 - no perfect synchrony?
 - tolerate faults?

Previously: String Matching.

- Given a text T[0..n) and a pattern P[0..m), determine if/where T contains P
- Focus on *one shot* complexity:
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A Variation. The text *T* is *fixed*, but we may wish to search *T* for many different (initially) unknown patterns $P_1, P_2, ...$

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An Alternative Approach. *Preprocess* the text *T* to make the searches more efficient

- Pay for preprocessing upfront
- Each query can be *much* more efficient.

Example Problem. Given a text *T* of *words*, implement an *index* of the occurrences of that word.

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Question. How can we implement such a map *efficiently*?

The Trie Data Structure

Idea. Store words in a tree

- · Each leaf represents a possible word in the text
- Each internal node represents *prefix* of a word in the text
 - path from root to leaf stores letters in the leaf word
- Append a terminating character to each word to make the tree a **prefix tree**
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Example: {aa\$, aaab\$, abaab\$, abb\$, abbab\$, bba\$, bbbb\$}



Question. Given a pattern *P* and a **trie** for the text *T*, how do we determine if *T* contains the pattern *P*?



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PollEverywhere

What is the running time of searching a trie?



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Remarkable fact. The time to search a trie depends only on the length of *P*, not the size of *T*!

• Also: the trie can be computed efficiently from *T* (in *O*(*n*) time).

Compact Tries

Observation. Tries are potentially wasteful!

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Idea. Compress paths without branches!

- Replace a path of **unary** (single-child) nodes with a single edge
- Label edge with the *first* character of the corresponding path
- Label each *vertex* with the index of the next character

Words, Trie, Compact Trie

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0

a

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abaah

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\$

Example. Search for ababb.

bbbb

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bb

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abh

abbab

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Useful feature. If a compact trie stores ℓ words, then it has at most

- $\ell 1$ internal nodes as well.
 - The size of the trie is proportional to the number of words it stores!
 - Fact (to prove). If a tree *T* has ℓ leaves and every internal node has at least two children, then *T* has at most $2\ell 1$ vertices.

Trie Discussion

Advantages of tries:

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 - DNA/RNA sequences
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We need new ideas!!

So Far.

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An observation. P[i, i+1), P[i, i+2), P[i, i+3),... can all just be checked against P[i, n)

Definition. Given a text T[0..n) the **suffix tree** \mathcal{T} of *T* is formed by:

- take the compact trie of all suffixes of T\$ (i.e., all $T_i = T[i..n)$ \$)
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- **Example.** T = banana

```
banana$
anana$
nana$
ana$
na$
a$
```



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PollEverywhere

Given T[0..n), what is the total size of the associated suffix tree \mathcal{T} ?



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For now. Take it as given that \mathcal{T} can be computed in O(n) time.

Suffix Tree Applications

Application 1: String Matching

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Observation. *P* occurs in $T \iff P$ is a prefix of a suffix of *T*.

- \mathcal{T} stores (references to) all suffixes in T
- To search for *P*, try follow a path with label *P* until
 - 1. we get stuck
 - internal node without next character
 - mismatch along an edge
 - 2. we reach end of pattern P
 - all descendent leaves contain P!
 - 3. reach a leaf ℓ with part of *P* left (no match)

Bananas. T = b a n a n a b a n

Human readible suffix tree:



Bananas. T = b a n a n a b a n

Human readible suffix tree:



Note. Operations on "human readable" tree can be simulated in true suffix tree.

· each internal node stores pointer to left-most descendant index

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Bananas. T = b a n a n a b a n

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Search P = baa

Bananas. T = b a n a n a b a n

Human readible suffix tree:



Search P =ana

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Search P = ba

String Matching Discussion

Using Suffix Trees

- Pre-process a text *T*[0..*n*) in *O*(*n*) time *once*
- Search for *P*[0..*m*) in time *O*(*m*) time

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Comparison. If *T* is large and static, and we expect to perform many searches, the suffix tree construction is *much* more efficient!

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Example. T = b a n a n a b a n **Repeated substrings** in the suffix tree?



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Example. T = b a n a n a b a n

Observation. Repeated substrings correspond to paths of internal nodes in \mathcal{T} .

- Longest repeated substring = longest path of internal nodes in \mathcal{T}
 - "longest path" includes weight for compressed edges
- Can be computed in *O*(*n*) time!
 - use "depth first search" strategy

Using suffix trees we can perform the following tasks efficiently:

- **1.** Longest Common Substring in time $O(n_1 + n_2 + \dots + n_k)$
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 - Output: the longest substring that is contained in all *T_i*

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4. Matching with Wildcards

- Input: text *T*[0..*n*), pattern *P*[0..*m*) with wildcards
 - wildcard character * matches a substring of any length
- Output: first appearance of *P* (with wildcard matches)

Conclusion

Suffix trees are amazing data structures!

- Tons of applications
- Surprising theoretical results

Next time. Constructing suffix trees efficiently

Scratch Notes