String comparison by transposition networks

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(joint work with Alexander Tiskin)

Strings, substrings and subsequences...

Find pattern **abab** occurring as substring in text bbbabababba \Rightarrow

bbb**abab**abba bbbab**abab**ba

Possible in O(n) time (Automata, Boyer Moore, Knuth Morris Pratt)

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Exact string comparison: Hamming distance

Count mismatches:

dist(bbbabababba, abbbbabaaba) = 3

Approximate string search

One way: subsequence matching Find pattern **abab** in text bbbabababba as a subsequence \Rightarrow

> bbb**aba**ba<mark>b</mark>ba bbb<mark>ab</mark>ab<mark>ab</mark>ba

Possible in O(n) time by constructing automata.

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> bbb**aba**ba**b**ba bbb**ab**ab**ab**ba

> > . . .

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Approximate comparison: string alignment

Align the maximum number of letters, preserving order:

abbabbbabbaba bbabaabbba

The aligned letters form the longest common subsequence (LCS); length of this sequence: LLCS.

dist(x,y) = m + n - 2 LLCS(x,y)

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The aligned letters form the longest common subsequence (LCS); length of this sequence: LLCS.

$$dist(x,y) = m + n - 2 LLCS(x,y)$$

Approximate comparison

LCS distance: Minimizes number of insertions/deletions to get from string x to string y.

Extensions we don't (directly) consider here:

- Edit distance: minimize the number of insertions, deletions, and exchange operations to get from one string to the other.
- Weighted case: assign weights to each operation on each pair of characters.

The LCS Problem

Complexities of classical solutions (input strings of length n, r matches, d dominant matches):

- Dynamic programming (Wagner & Fischer, '74) : O(n²)
- ⇒ Using "Four Russians" technique (Masek & Paterson, '80) : $O(n^2/\log n)$
 - Dominant match based (Hunt & Szymanski, '77) : O((r + n) log n) (Apostolico & Guerra, '87) : O(m log n + d log(mn/d))

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Seaweeds and networks...

What is semi-local string comparison?

Semi-local comparison: compute substring-string *highest-score matrix*

$$A(i,j) = LLCS(x_i x_{i+1} \dots x_j, y)$$

and simultaneously all string-substring, prefix-suffix and suffix-prefix LLCS.

Why compare semi-locally?

- Two N \times N highest-score matrices can be (min,+) multiplied in $O(N^{1.5})$
- ⇒ Semi-local comparison is useful for obtaining efficient parallel LCS algorithms
 - Semi-local comparison has non-trivial algorithmic applications itself.
 - One step closer to fully local comparison (substring vs. substring)

How to do it:

Use the Seaweed Algorithm, which runs in $O(n^2)$

Start with *extended alignment-dag*:



- We trace seaweeds through cells
- Two seaweeds cross at most once
- We are interested in start and end points of seaweeds



Querying the LCS distance by counting seaweeds:



Querying the LCS distance by counting seaweeds:



Comparison networks

Comparison network:

- n wires connected by arbitrary number of comparators
- Comparators have two inputs and two outputs.
 - \Rightarrow return larger value the predefined output
 - \Rightarrow return smaller value at the other output.
- Traditional method for studying oblivious sorting algorithms

Transposition network: all comparators only connect adjacent wires.

Merging using a transposition network

Example (Merging two 4 element sorted sequences)



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[Munter, U.S. Patent 5,216,420 '93] We call this a DIAMOND(4,4) network.

Observation

Every mismatch cell in the alignment dag behaves as a comparator.

Therefore

... we can define a transposition network to solve LCS problem.



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Definition (The LCSNET(x, y) network)



Remove all comparators in DIAMOND(|x|, |y|) which correspond to matches.

Definition (The LCSNET(x, y) network)



Initialize network with |x| ones and |y| zeros.

Definition (The LCSNET(x,y) network)



Count the number of ones which reach the right hand side \Rightarrow LCS distance.

An application: Sparse string comparison...

Kinds of sparseness

Few match cells in the alignment dag

Extreme case: comparison of permutation strings

Strings very similar (LCS is long / edit distance is short)

Not necessarily correlated with number of matches

Strings very dissimilar (LCS is short) Correlated with number of matches, but still different measure

Obtaining all matches

- We want to work match by match...
- \Rightarrow How to obtain list of matches in less than $O(n^2)$ time?
- For small alphabets ($|\Sigma| < n$): in $O(n \log |\Sigma|)$ by counting character frequencies
- For large alphabets: in O(n log n) by sorting one of the input strings and binary search

(if characters can only be tested for equality, $\Omega(n^2)$ is a lower bound)

Classical approach for sparse comparison

Trace antichains and their contours:



(a) input strings and alignment (b) prefix-prefix LCS lengths dag and contours



















More about cells

Not all cells in the 0-1 transposition network need to be evaluated:



Parameterized LCS computation

We obtain a simple algorithm for computing the LCS in O(p(n-p)) by tracing zeros and ones (which is equivalent to the best known result).

Parameterized LCS computation

We can also obtain an algorithm for semi-local comparison which runs in O(np):



Summary

- Transposition networks provide unified view on different LCS/semi-local comparison algorithms
- Transposition networks allow to derive parameterized algorithms for global LCS more easily than existing approaches.
- We have new parameterized algorithms for sparse semi-local string comparison running in O(np + preproc(n)) and $O(n\sqrt{r} + preproc(n))$

Outlook

Some further work:

- Generalize to other types of distances (edit distance).
- Apply techniques to LCS with non-overlapping inversions
- Another interesting problem: string comparison in streaming models (i.e. we can only read strings once/predefined number of times)

Thanks!

... any guestions?

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