String indexing in the Word RAM model, part 3

Paweł Gawrychowski

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What we exactly mean by "defined" depends on the exact version. The most common are the following two:

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The blocks are described by pairs (in LZW) or triples (in LZ):

...ababbabaaabbabaabaabbabaaa... ...,a,b,(1,2,b),(1,4,a),(1,1,a),(4,8,b),(11,4,b),(10,2,a),...

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$$\int_{p=aaab} p=aaab$$

We want to store repetitive texts (say, genomic databases) in compressed form, but such that we can search them quickly.

In other words, given a text, build a small structure which allows fast pattern matching.

Pattern matching?

Given *p*[1..*m*] we want to find where it occurs **exactly** in text *t*[1..*n*]. We might want the first occurrence, or all of them, or just a few...

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Problem, more precisely

We are asked to build a self-index for a string t[1..n] whose LZ77 parse consists of *z* phrases.

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The number of those phrases is believed to be the **right** measure of how repetitive the text is.

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Straight-line program, or grammar representation

Simply a context-free grammar with **exactly** one production per nonterminal.

Rytter 2003, Charikar et al. 2005

A LZ77 parse consisting of *z* phrases can be converted to a grammar consisting of $g = O(z \log n)$ words. The grammar is AVL-balanced (Rytter) or weight-balanced (Charikar et al.).

Weight-balanced means that for each production $A \rightarrow BC$ we have that $|B| \approx |C|$.

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Framework (of Navarro)



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Observation (by Kärkkäinen and Ukkonen?)

If the pattern occurs in the text, there is at least one primary occurrence.

Assuming we have all primary occurrences, all secondary occurrences can be found via 2-sided 2D range reporting.

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Secondary occurrence

An occurrence is secondary iff it is completely contained in some phrase.

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Idea, continued



To find all primary occurrences of p[1..m], for each $1 \le i \le m$, we

- search for p[i + 1..m] in the compacted trie of the suffixes starting at phrase boundaries,
- search for (p[1..i])^R in the compacted trie of the reversed phrases,
- check the results via random access,
- use range reporting to find all boundaries preceded by p[1..i] and followed by p[i + 1..m].

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Bookmarking

Because we know that we will extract characters from the phrase boundaries, we can replace $O(\log n + \ell)$ with the following bound:

Lemma

Given a balanced SLP for S with g rules and integers b and L, we can store $2 \log g + \mathcal{O}(\log L)$ bits such that later, given $\ell \leq L$, we can extract $t[b - \ell..b + \ell]$ in $\mathcal{O}(\log L + \ell)$ time.

Corollary

Given b, we can store $\mathcal{O}(\log^* z)$ words such that, given any ℓ , we can extract $t[b - \ell .. b + \ell]$ in $\mathcal{O}(\ell)$ time.

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Space bounds (in words)

Patricia trees $\mathcal{O}(z)$ bookmarks $\mathcal{O}(z \log^* z)$ 4-sided 2D range reporting $\mathcal{O}(z \log \log z)$ 2-sided 2D range reporting $\mathcal{O}(z)$

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Time bounds

searching in compacted tries (with perfect hashing if necessary) extracting from bookmarks 4-sided 2D range reporting 2-sided 2D range reporting $\mathcal{O}(m^2)$

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 $\mathcal{O}(m \log \log n)$

 $\mathcal{O}(occ \log \log n)$

 $\mathcal{O}(m^2 + (m + occ) \log \log n)$

A simple trick to remove the $m \log \log n$:

For each node of the compacted trie (of the prefixes), store a 1D range reporting structure with the pre-orders in the other compacted trie:

Alstrup, Brodal, Rauhe STOC 2001

1D range reporting on z points can be solved in O(z) space and optimal O(1 + occ) query time.

If $m \le \log \log n$ then use the 1D range reporting structure, otherwise m^2 subsumes the $m \log \log n$ anyway.

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Final result

Theorem

Given a balanced SLP for a string t[1..n] whose LZ77 parse consists of *z* phrases, we can add $\mathcal{O}(z \log \log z)$ words such that, given a pattern p[1..m], we can find all occ occurrences of *p* in $\mathcal{O}(m^2 + occ \log \log n)$ time.

Can we decrease m^2 ?

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A bit technical, but let's try!

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Checking if two strings are equal can be done by comparing their fingerprints:

Karp-Rabin-style fingerprints

$$\phi(\boldsymbol{s}) = \sum_{k=1}^{|\boldsymbol{s}|} \boldsymbol{S}[k] \sigma^{|\boldsymbol{s}|-k} \bmod p$$

Lemma

For a prime p and $r \in \{1, 2, ..., p-1\}$ chosen uniformly at random, the probability that $\phi_r(s) = \phi_r(s')$ even though $s \neq s'$ is at most $\frac{|s|}{p-1}$.

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z-fast tries

We want to preprocess a compacted trie T on n nodes for navigating with a query string x. In this application, it is enough to find the unique (implicit or explicit) node of T that corresponds to the whole x, if such a node exists, and otherwise return any node. However, the procedure will in fact do a bit more.

2-fattest number

The 2-fattest number in a nonempty interval of positive integers is the number in the interval whose binary representation has the highest number of trailing zeros

For every edge of T, we choose the implicit node on the edge whose string depth is the 2-fattest number in the corresponding range, and make it explicit.

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Search in a z-fast trie

This can be done in only $\mathcal{O}(\log |x|)$ iterations:

```
Algorithm 1 Querying the probabilistic z-fast trie
(represented by the function T).
  input x \in u
  i \leftarrow \lceil \log w \rceil - 1
  \ell, r \leftarrow 0, w
  while r - \ell > 1 do
      if \exists b such that 2^i b \in (\ell \dots r) then
                       \{2^i b \text{ is the 2-fattest number in } (\ell \dots r)\}
         q \leftarrow \text{prefix of } x \text{ of length } 2^i b
         \langle q, s \rangle \leftarrow T(q)
         if g \leq |x| and s is the signature of the prefix of
         x of length g then
                                  {Move from (\ell \dots r) to (q \dots r)}
            \ell \leftarrow q
         else
            r \leftarrow 2^i b
                        {Move from (\ell \dots r) to (\ell \dots 2^i b)}
         end if
      end if
      i \leftarrow i - 1
  end while
  return \ell
```

For a given suffix p[i..m], this allows us to find the unique (implicit or explicit) node of the compacted trie.

Fingerprinting

Given a balanced SLP of size g, we can store $\mathcal{O}(g)$ words of extra information such that we can compute the fingerprint of any substring in $\mathcal{O}(\log n)$ time.

Bookmarked fingerprinting

Given a balanced SLP of size g and an integer b, we can store $\mathcal{O}(\log \log n)$ words of extra information such that later, given ℓ , we can compute the fingerprint of any $t[b..b + \ell)$ in $\mathcal{O}(\log \ell)$ time.

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We now have queries in $\mathcal{O}(m \log m + (m + occ) \log \log n)$ time and $\mathcal{O}(z \log n)$ space. How to remove $m \log \log n$?

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Bille, Cording, Gørtz, Sach, Vildhøj, Vind WADS 2013

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What's next?

Let r be the number of runs in the BWT of the text.

Gagie, Navarro, Prezza JACM 2020

An index taking $\mathcal{O}(r \log(n/r))$ words and generating all *occ* occurrences in $\mathcal{O}(m + occ)$ time.

Let $\delta = \max_{\ell=1}^{n} d_{\ell}/\ell$, where d_{ℓ} is the number of distinct length- ℓ substrings of the text.

Kempa and Kociumaka FOCS 2023

An index taking $\mathcal{O}(\delta \log \frac{n \log \sigma}{\delta \log n})$ words and allowing suffix array and inverse suffix array queries in $\mathcal{O}(\log^{4+\epsilon} n)$ time.

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