

On the Benefit of Merging Suffix Array Intervals for Parallel Pattern Matching

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71. Workshop über Algorithmen und Komplexität

- Σ is the alphabet with $|\Sigma| = \sigma$
- $\$ \notin \Sigma$ and $\forall \alpha \in \Sigma : \$ <_{\text{lex}} \alpha$
- $T \in \Sigma^* \cup \{\$\}$ and $P \in \Sigma^*$
- $|T| = n$ and $|P| = m$
- p is the number of processors

Pattern Matching

Given a text T of length n and a pattern P of length m , find all occurrences of P in T .

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$T = \text{b} \text{anana} \$$

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Pattern Matching

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$P_1 = \text{b}$ and $P_2 = \text{a}$

Sequential Times

Type	Query Time	Idea
exact	$\mathcal{O}(m)$	Suffix Tree
k -errors	$\mathcal{O}(m^k \sigma^k \max(k, \lg \lg n) + occ)$	[Lam et al., 2007]

Prefix and Suffix

$P_i = T[1..i]$ is the i -th prefix of T for all $i \in [1, n]$

$S_i = T[i..n]$ is the i -th suffix of T for all $i \in [1, n]$

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i	1	2	3	4	5	6	7
S_i	banana\$	anana\$	nana\$	ana\$	na\$	a\$	\$

Suffix Array of T

The SA is a permutation of $[1, n]$ such that for all $i \in [1, n - 1]$:

$$T[SA[i]..n] <_{\text{lex}} T[SA[i + 1]..n]$$

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	1	2	3	4	5	6	7
SA[i]	7	6	4	2	1	5	3
	\$	a	a	a	b	n	n
		\$	n	n	a	a	a
			a	a	n	\$	n
			\$	n	a		a
				a	n		\$
				\$	a		
					\$		

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Suffix Array Interval (SAI) of P

$$i \in \mathcal{I}(P) \iff T[SA[i]..SA[i] + |P| - 1] = P$$

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$$\mathcal{I}(a) = [2, 4]$$

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\$		a	a	a	b	n	n
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a				a	n	\$	n
\$					n	a	a
a						a	n
\$							a
\$							
\$							

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	1	2	3	4	5	6	7
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	\$	a	a	a	b	n	n
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Suffix Array Interval (SAI) of P

$$i \in \mathcal{I}(P) \iff T[SA[i]..SA[i] + |P| - 1] = P$$

Inverse Suffix Array of T

The SA^{-1} is a permutation of $[1, n]$ such that for all $i \in [1, n]$:

$$SA^{-1}[SA[i]] = i$$

The Inverse Suffix Array

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$$\mathcal{I}(a) = [2, 4]$$

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	1	2	3	4	5	6	7
$SA[i]$	7	6	4	2	1	5	3
$SA^{-1}[i]$	5	4	7	3	6	2	1
	\$	a	a	a	b	n	n
		\$	n	n	a	a	a
			a	a	n	\$	n
			\$	n	a		a
				a	n		\$
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	1	2	3	4	5	6	7
$SA[i]$	7	6	4	2	1	5	3
$\Psi^1[i]$	-	1	6	7	4	2	3
	\$	a	a	a	b	n	n
		\$	n	n	a	a	a
			a	a	n	\$	n
			\$	n	a		a
				a	n		\$
				\$	a		
					\$		

Find the rest of the suffix

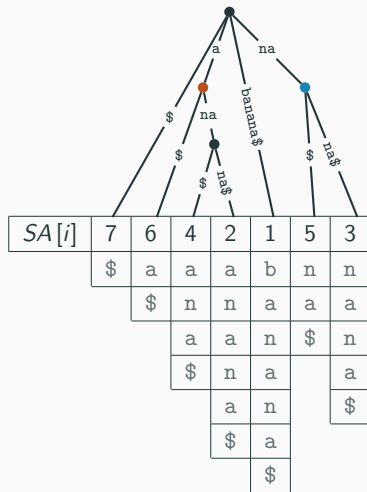
$$\Psi^k[i] = SA^{-1}[SA[i] + k]$$

Tree above the Suffix Array

- Nodes cover *relevant SAs*

$$I(a) = [2, 4]$$

$$I(n) = [6, 7]$$



Suffix Array Interval Merging

The Idea

Find occurrences of subpatterns and merge suffix array intervals

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The Problem

How to find the interval gained by merging two suffix array intervals?

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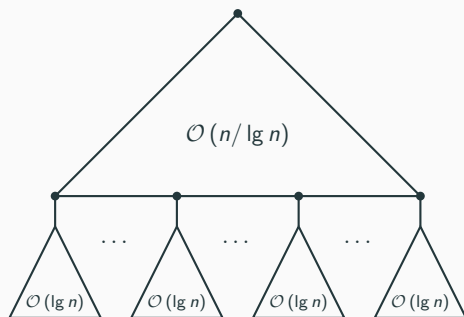
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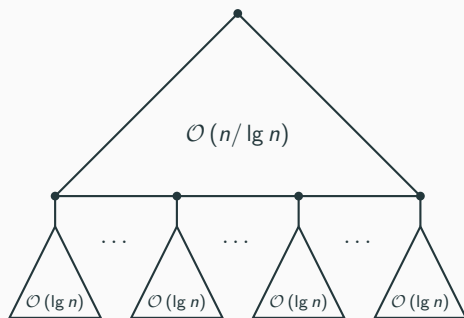
How to find the interval gained by merging two suffix array intervals?

Paper	Running Time	Idea
[Huynh et al., 2006]	$\mathcal{O}(\lg n)$	Binary Search
[This talk]	$\mathcal{O}(\lg \lg n)$	Extending [Lam et al., 2007], Sampling Ψ in y -fast trie
	$\mathcal{O}(\lg_p \lg n)$	Parallel Binary Search



y-Fast Trie [Willard, 1983]

- Each leaf stores $\mathcal{O}(\lg n)$ elements in a binary search tree
- x-fast trie for $\mathcal{O}(n/\lg n)$ elements
- Prefixes of elements in $\mathcal{O}(\lg n)$ hash tables



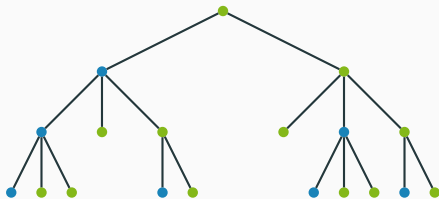
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FIND, PREDECESSOR and SUCCESSOR in $\mathcal{O}(\lg \lg n) \dots$

- ... expected time **or**
- ... deterministic time with $\mathcal{O}(n \lg \lg n)$ construction time.

Heavy Path Decomposition

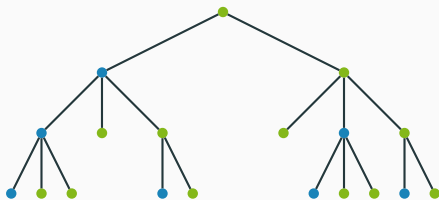


Nodes are

Heavy if they are in the largest subtree

Light otherwise (or if they are the root)

Heavy Path Decomposition



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Sample ψ for each **light** node

Given two SAs $\mathcal{I}(\alpha)$ and $\mathcal{I}(\beta)$

- Find all $i \in \mathcal{I}(\alpha) : \Psi^{|\alpha|} [i] \in \mathcal{I}(\beta)$
- $\Psi^{|\alpha|} [i]$ is monotonically increasing for all $i \in \mathcal{I}(\alpha)$

Given two SAls $\mathcal{I}(\alpha)$ and $\mathcal{I}(\beta)$

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Sampling for **Light Nodes** v of $\mathcal{I}(\alpha)$ in y -fast trie

$$\Gamma(v) := \{(\Psi^{|\alpha|}[i], i) : i \equiv 1 \pmod{\lg^2 n} \wedge i \in \mathcal{I}(\alpha)\}$$



Given two SA/Is $\mathcal{I}(\alpha)$ and $\mathcal{I}(\beta)$

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Merging SA/s – Light nodes

Let v be the **light** node of $\mathcal{I}(\alpha)$ and $\mathcal{I}(\beta) = [b_\beta, e_\beta]$

- If $\Gamma(v) = \emptyset \rightarrow$ Binary search on $< \lg^2 n$ elements



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- Find $j_l, j_r : b_\beta \leq \Psi^{|\alpha|}[j_l]$ and $\Psi^{|\alpha|}[j_r] \leq e_\beta$
- Extend j_l, j_r using binary search on $< \lg^2 n$ elements



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Find $k_l, k_r : \Psi^{|\alpha|}[k_l] \leq b_\beta$ **and** $e_\beta \leq \Psi^{|\alpha|}[k_r]$

- Shrink k_l, k_r using binary search on $< \lg^2 n$ elements



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Similar idea for **heavy** nodes

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Lemma

We can merge two SAIs in $\mathcal{O}(\lg \lg n)$ time.

What are we doing to merge two *SA*'s

Query *y*-fast tries **and** binary search

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Parallelize these queries

- Binary search requires $\mathcal{O}(\lg_p n)$ parallel time [Snir 1985]
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- Static *y*-fast trie \rightarrow arrays instead of binary search trees

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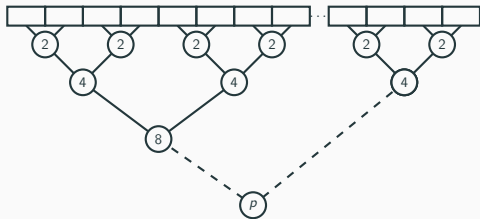
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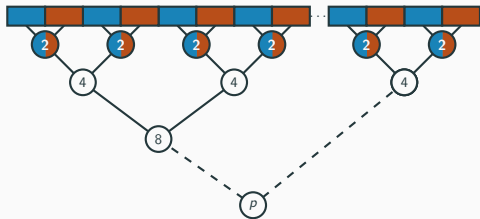
Parallel Exact Pattern Matching

- $P = P_1P_2 \dots P_p$ with $|P_i| = m/p$
- Compute $\mathcal{I}(P_i)$ in $\mathcal{O}(m/p)$ time
- Merge $SAIs$ in $\mathcal{O}(\lg_p \lg n)$ time



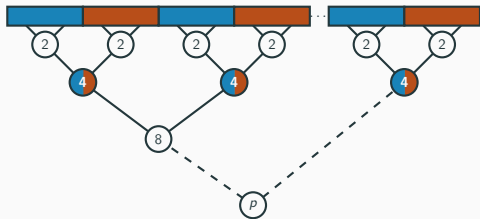
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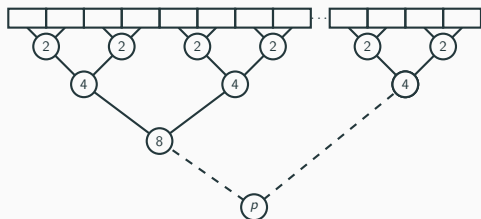
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In the k -th Step

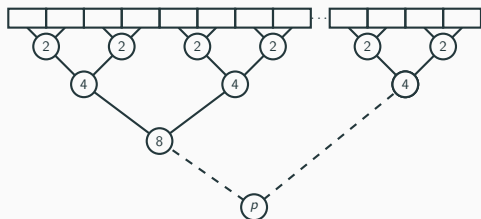
$p/2^k$ SAIs $\rightarrow 2^k$ processors

Number of Steps

There are $\lg_p \lg n$ merge steps

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Theorem

Parallel exact pattern matching requires $\mathcal{O}(m/p + \lg \lg p \lg \lg n)$ time.

The k -Difference and k -Mismatch Problem

Given a text T of length n and a pattern P of length m ...

k -Difference Problem

... find all occurrences of P' in T such that P can be transformed to P' using $\leq k$ INSERT, CHANGE and DELETE operations.

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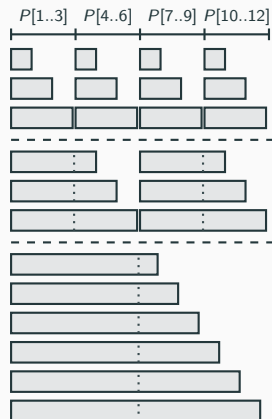
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k -Mismatch Problem

... find all occurrences of P' in T such that P can be transformed to P' using $\leq k$ CHANGE operations.

Compute *SA*/s of all prefixes and suffixes of P

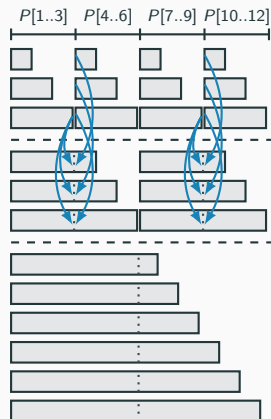
Preprocessing: $|P| = 12, p = 4$



Preprocessing

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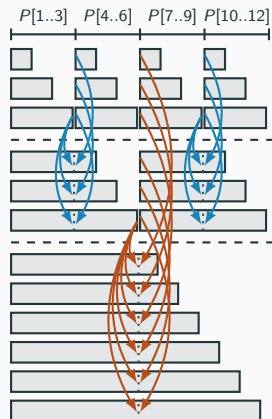
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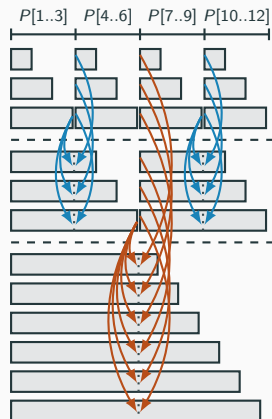


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- $p/2^k$ left *SAI*s
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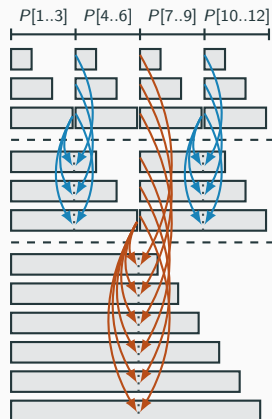
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Cost of Merging

- There are $\lg n$ merge steps
- Merging in $\mathcal{O}(\lg_p \lg n)$ time



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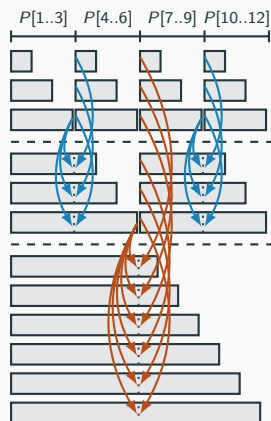
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Lemma

The preprocessing requires $\mathcal{O}(m/p \lg p \lg \lg n)$ time.



Introducing the Error (**Insert**, **Change** or **Delete**)

- $\mathcal{I}(P[1..i])$ and $\mathcal{I}(P[i..n])$ are known
- What is an error at position j

P

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Solving the 1-Difference and 1-Mismatch Problem

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Theorem

Approximate parallel pattern matching with ≤ 1 error can be solved in $\mathcal{O}(\sigma m/p \cdot \lg \lg n + occ)$ time.

Solving the k -Difference and k -Mismatch Problem

Quite similar to $k = 1$

- The same preprocessing
- Introduce $\leq k$ errors by merging SA 's
- Use configurations of positions and parallelize those



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Problem – Report Occurrence Multiple Times

The Problem: $T = \text{aaa}\$$ and $P = \text{aba}$ and one error

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Both P' and P'' occur at position 1 in T

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The Solution

- Report only if found with smallest distance [Huynh et al., 2006]
- Can be parallelized

Things we did

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Thank You