# Sampled LCP Array 

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## Outline

## 1. Introduction

2. Irreducible LCP Algorithm
3. Sampled LCP Array
4. Conclusions

## Suffix Trees

- Important data structures in string processing and bioinformatics.
- In practice: 10 - 20 bytes / character with 32-bit pointers (Kurtz 1999).
- For large data sets, we want something much smaller but still efficient.


## Space-Efficient Alternatives

## SA BWT

| 12 | I | SA | SA: $(\log n+\log \sigma)$ bits / char |
| :---: | :---: | :---: | :---: |
| 11 | P | I\$ |  |
| 8 | S | IPPI\$ |  |
| 5 | S | ISSIPPI\$ |  |
| 2 | M | ISSISSIPPI\$ |  |
| 1 | \$ | MISSISSIPPI | PI\$ |
| 10 | P | PI\$ |  |
| 9 | I | PPI\$ |  |
| 7 | S | SIPPI\$ |  |
| 4 | S | SISSIPPI\$ | functionality! |
| 6 | I | SSIPPI\$ |  |
| 3 | I | SSISSIPPI\$ |  |

## Compressed Suffix Trees

- Many proposals have three components:
- Compressed suffix array (CSA)
- Longest common prefix (LCP) array
- Tree topology
- In practice: Cánovas, Navarro (SEA 2010)


## Longest Common Prefix Array

| SA | BWT | LCP |  |
| ---: | :---: | :---: | :--- |
|  |  |  |  |
| 12 | I | 0 | \$ |
| 11 | P | 0 | IS |
| 8 | S | 1 | IPPIS |
| 5 | S | 1 | ISSIPPI\$ |
| 2 | M | 4 | ISSISSIPPI\$ |
| 1 | $\$$ | 0 | MISSISSIPPI\$ |
| 10 | P | 0 | PIS |
| 9 | I | 1 | PPI\$ |
| 7 | S | 0 | SIPPI\$ |
| 4 | S | 2 | SISSIPPI\$ |
| 6 | I | 1 | SSIPPI\$ |
| 3 | I | 3 | SSISSIPPI\$ |

## Longest Common Prefix Array

| SA | BWT | LCP |  |  |
| ---: | :---: | :---: | :--- | :--- |
|  |  |  |  |  |
| 12 | I | 0 | \$ |  |
| 11 | P | 0 | IS |  |
| 8 | S | 1 | IPPI\$ | Both preceded by 'S' |
| 5 | S | 1 | ISSIPPIS |  |
| 2 | M | 4 | ISSISSIPPI\$ |  |
| 1 | $\$$ | 0 | MISSISSIPPI\$ |  |
| 10 | P | 0 | PI\$ |  |
| 9 | I | 1 | PPIS |  |
| 7 | S | 0 | SIPPI\$ | Previous suffixes |
| 4 | S | 2 | SISSIPPI\$ | are also adjacent. |
| 6 | I | 1 | SSIPPI\$ |  |
| 3 | I | 3 | SSISSIPPI\$ |  |

## Longest Common Prefix Array

SA BWT LCP

| 12 | I | 0 | \$ |
| :---: | :---: | :---: | :---: |
| 11 | P | 0 | I \$ |
| 8 | S | 1 | IPPI\$ |
| 5 | S | 1 | ISSIPPI\$ |
| 2 | M | 4 | ISSISSIPPI\$ |
| 1 | \$ | 0 | MISSISSIPPI\$ |
| 10 | P | 0 | PIS |
| 9 | I | 1 | PPI\$ |
| 7 | S | 0 | SIPPI\$ |
| 4 | S | 2 | SISSIPPI\$ |
| 6 | I | 1 | SSIPPI\$ |
| 3 | I | 3 | SSISSIPPI\$ |

Suffix and left match preceded by same character.

## Permuted LCP Array

| SA | BWT | PLCP |  |
| :---: | :---: | :---: | :---: |
| 1 | \$ | 0 | MISSISSIPPI\$ |
| 2 | M | 4 | ISSISSIPPI\$ |
| 3 | I | 3 | SSISSIPPI\$ |
| 4 | S | 2 | SISSIPPI\$ |
| 5 | S | 1 | ISSIPPI\$ |
| 6 | I | 1 | SSIPPI\$ |
| 7 | S | 0 | SIPPI\$ |
| 8 | S | 1 | IPPI\$ |
| 9 | I | 1 | PPI\$ LCP[x] = PLCP[SA[x]] |
| 10 | P | 0 | PI\$ |
| 11 | P | 0 | I\$ |
| 12 | I | 0 | \$ |

## Compressing the LCP Array

- PLCP[i] + $2 i$ is strictly increasing, and can be represented as a bit vector of length $2 n$.
- Sadakane (2007)
- There are at most R runs of ones in the bit vector, so we can use run-length encoding.
- Fischer, Mäkinen, Navarro (2009)


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## Maximal and Minimal Values

| SA | BWT | PLCP |  | MAX | MIN |
| ---: | :---: | :---: | :--- | :---: | :---: |
|  |  |  |  |  |  |
| 1 | $\$$ | 0 | MISSISSIPPIS | X | X |
| 2 | M | 4 | ISSISSIPPI\$ | X |  |
| 3 | I | 3 | SSISSIPPI\$ |  |  |
| 4 | S | 2 | SISSIPPI\$ |  |  |
| 5 | S | 1 | ISSIPPI\$ | X | X |
| 6 | I | 1 | SSIPPIS | X | X |
| 7 | S | 0 | SIPPI\$ | X | X |
| 8 | S | 1 | IPPIS | X | X |
| 9 | I | 1 | PPI\$ | X | X |
| 10 | P | 0 | PI\$ | X | X |
| 11 | P | 0 | IS | X | X |

## Irreducible LCP Algorithm

- Kärkkäinen, Manzini, Puglisi (CPM 2009)
- Find and compute maximal values.
- Their sum is at most $2 \mathrm{n} \log \mathrm{n}$.
- Other values: $\operatorname{PLCP}[i+1]=\operatorname{PLCP}[i]-1$
- PLCP construction: $O(n$ log $n$ ) time, negligible working space.
- Requires the text in memory and SA on disk.
- How to use a CSA instead?


## Compressed Suffix Arrays

- Function $\Psi: \operatorname{SA}[\Psi(x)]=\operatorname{SA}[x]+1$
- Scan CSA in text order: Start from (i, x) $=\left(i, S A^{-1}[i]\right)$ and iterate with $(i+1, \Psi(x))$.
- Array $\mathrm{C}: \mathrm{C}[\mathrm{c}]$ is the number of occurrences of characters in $\{0, \ldots, c-1\}$.
- Function $\Psi$ is strictly increasing in every

$$
\Psi_{C}=[C[c-1]+1, C[c]] .
$$

- SA samples: Compute $S A[x]=i$ and $S A^{-1}[i]=x$ when $i$ is a multiple of parameter $d$.
- $S A[x]=S A\left[\Psi^{k}(x)\right]-k$.


## Modified Algorithm

- Scan the CSA in text order, starting from (1, $\left.\mathrm{SA}^{-1}[1]\right)$.
- If PLCP[i] is maximal, compute it by scanning the CSA from ( $\mathrm{i}, \mathrm{x}$ ) and its left match ( $\mathrm{j}, \mathrm{x}-1$ ).
- Otherwise PLCP[i] = PLCP[i-1] - 1.
- Time complexity is $O\left(t_{\psi} n \log n\right)$, and working space is still negligible.


## Identifying Maximal Values

- Suffix i of rank $x$ and its left match $j$ of rank $x-1$.
- PLCP[i] is non-maximal iff BWT[x] = BWT[ $x-1]$.
- PLCP[i] is non-maximal iff the ranks of suffixes $\mathrm{i}-1$ and $j-1$ are in the same $\Psi_{c}$.
- Let $y$ be the rank of suffix $i-1$. PLCP[i] is nonmaximal iff $y-1$ and $y$ are in the same $\Psi_{c}$, and $\Psi(y-1)=x-1$.


## In Practice

| Name | Size | Time | MB / s |
| :--- | ---: | ---: | ---: |
| English | 400 MB | 1688 s | 0.24 |
| Fiwiki | 400 MB | 327 s | 1.22 |
| DNA | 385 MB | 3475 s | 0.11 |
| Yeast | 409 MB | 576 s | 0.71 |

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## Using PLCP with a CSA

- We want to retrieve $\operatorname{LCP}[x]=\operatorname{PLCP}[S A[x]]$.
- To get $\operatorname{SA}[x]$, we iterate $\Psi^{k}(x)$ for $k=0,1, \ldots$, until we find a sampled $\operatorname{SA}\left[\Psi^{k}(x)\right]$ value.
- Then LCP $[x]=$ PLCP $\left[S A\left[\Psi^{k}(x)\right]-k\right]$.
- Time complexity is $\mathrm{O}\left(\mathrm{d}_{4}\right)$.
- Typical values of d are 16 to 64 for regular texts, and 128 to 512 for highly repetitive texts.
- $t_{\psi}$ is roughly $1 \mu \mathrm{~s}$ in practice.


## A Faster Approach

- Cánovas, Navarro (SEA 2010)
- LCP values are usually small, so we can compress the LCP array directly.
- For regular texts, we get $6-8$ bits / character.
- PLCP is at most 2 bits / character.
- $\operatorname{LCP}[x]$ can be retrieved in less than $1 \mu \mathrm{~s}$.


## Sampled LCP Array (1 / 2)

- $\operatorname{PLCP}[i]=\operatorname{PLCP}[i+1]+1$, if $i$ is non-minimal.
- PLCP[i] = PLCP $[i+k]+k$, if $i+k$ is the next minimal value.


## Maximal and Minimal Values

| SA | BWT | PLCP |  | MAX | MIN |
| ---: | :---: | :---: | :--- | :---: | :---: |
|  |  |  |  |  |  |
| 1 | $\$$ | 0 | MISSISSIPPIS | X | X |
| 2 | M | 4 | ISSISSIPPI\$ | X |  |
| 3 | I | 3 | SSISSIPPI\$ |  |  |
| 4 | S | 2 | SISSIPPI\$ |  |  |
| 5 | S | 1 | ISSIPPI\$ | X | X |
| 6 | I | 1 | SSIPPIS | X | X |
| 7 | S | 0 | SIPPI\$ | X | X |
| 8 | S | 1 | IPPIS | X | X |
| 9 | I | 1 | PPI\$ | X | X |
| 10 | P | 0 | PI\$ | X | X |
| 11 | P | 0 | IS | X | X |

## Sampled LCP Array (1 / 2)

- $\operatorname{PLCP}[i]=\operatorname{PLCP}[i+1]+1$, if $i$ is non-minimal.
- PLCP[i] = PLCP $[i+k]+k$, if $i+k$ is the next minimal value.
- We store the R minimal PLCP values in SA order.
- Additional samples may be needed for performance.
- To get LCP[x], we iterate $\Psi^{k}(x)$ for $k=0,1, \ldots$, until we find a sampled $\operatorname{LCP}\left[\Psi^{k}(x)\right]$ value.
- Then $\operatorname{LCP}[x]=\operatorname{LCP}\left[\Psi^{k}(x)\right]+k$.


## Sampled LCP Array (2 / 2)

- The size of the minimal samples scales with R.
- Somewhat larger in practice than a run-length encoded PLCP.
- In regular texts, $20-40 \%$ of the LCP values are minimal.
- LCP values can be retrieved several times faster than by using a PLCP array.
- On highly repetitive texts, the performance is determined by the number of extra samples.


## Experiments: English



## Experiments: Fiwiki



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## Conclusions

- We can construct the (P)LCP array directly from a CSA with little extra working space.
- Construction speed is similar to direct CSA construction.
- The LCP array can be sampled in a similar way as the suffix array.
- On regular texts, the sampled LCP array offers better time/space trade-offs than the PLCP array.

