

Compressed Suffix Trees in Practice

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Outline

1 Introduction

- Basic data structures
- The suffix tree

2 CST design

- NAV (tree topology and navigation)
- CSA (lexicographic information)
- LCP (longest common prefixes)

3 CST in practice

- The *sds/* library

Succinct data structures (1)

Data structure D

representation of an
object X

+

operations on X

Example: Rank-bit-vector

bit vector b of length n

$(0, 1, 0, 1, 1, 0, 1, 1)$

$(0, 0, 1, 1, 2, 3, 3, 4)$

in n bits space

access $b[i]$ in $\mathcal{O}(1)$ time
 + $rank(i) = \sum_{j=0}^{i-1} b[j]$ in $\mathcal{O}(n)$ time

Succinct data structure D

Space of D is close the information theoretic lower bound to represent X , while operations can still be performed efficient.

Succinct data structures (2)

Can succinct data structures replace classic uncompressed data structures *in practice*?

- Less memory \Rightarrow fewer CPU cycles !?
- Less memory \Rightarrow less costs !?

Problems:

- in theory
 - develop succinct data structures
- in practice
 - constants in $\mathcal{O}(1)$ -time terms are large
 - $o(n)$ -space term is not negligible
 - complex data structures are hard to implement

Succinct data structures (2)

Can succinct data structures replace classic uncompressed data structures *in practice*?

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- Less memory \Rightarrow less costs !?

Instance name	main memory	price per hour
Micro	613.0 MB	0.02 US\$
High-Memory Quadruple Extra Large	68.4 GB	2.00 US\$

Pricing of Amazons Elastic Cloud Computing (EC2) service in July 2011.

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The classic index data structure: The suffix tree (ST)

Let T be a text of length n over alphabet Σ of size σ .

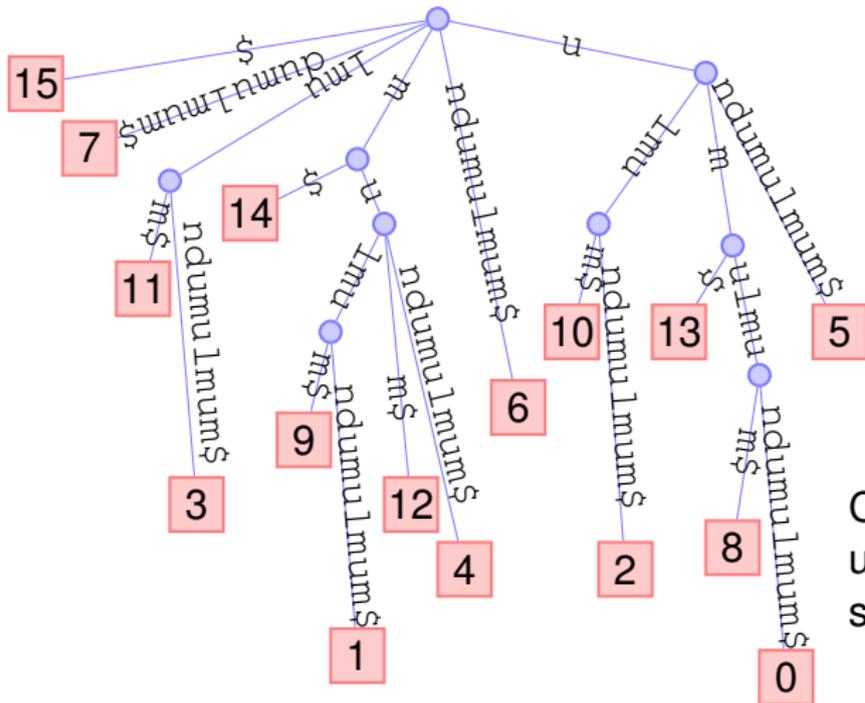
Suffix tree

- index data structure for T (construction $\mathcal{O}(n)$)
- can be used to solve many problems in optimal time complexity
 - bioinformatics
 - data compression
- uses $\mathcal{O}(n \log n)$ bits!
In practice (ASCII-alphabet) ≥ 17 times the size of T

Can not handle „The Attack of Massive Data”

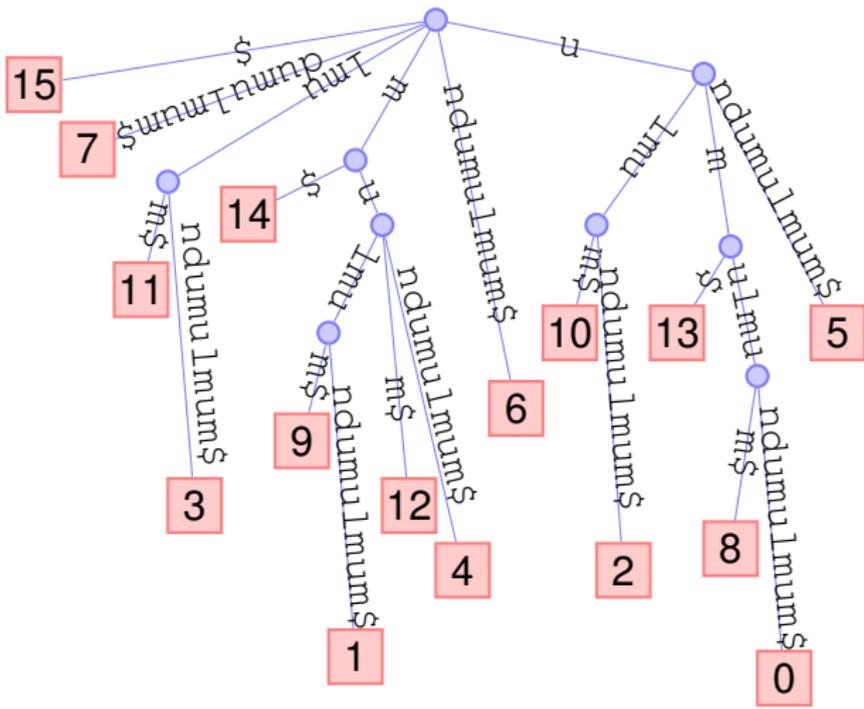
- DNA sequencing data (NGS)
- ...

Example: ST of $T = \text{umulumdumulum}\$$


 $n = 16$
 $\Sigma = \{\$, d, l, m, n, u\}$
 $\sigma = 6$

Classic implementation
uses pointers each of
size 4 or 8 bytes!

Example: ST of $T = \text{umulumdumulum}\$$



Operations

- root()*
- is_leaf(v)*
- parent(v)*
- degree(v)*
- child(v, c)*
- select_child(v, i)*
- depth(v)*
- edge(v, d)*
- lca(v, w)*
- sl(v)*
- wl(v, c)*

CSTs

Goal of a CST implementation

Replace fastest uncompressed ST implementations in different scenarios

- (a) both fit in RAM and we measure time
- (b) both fit in RAM and we measure resource costs
- (c) only CST fits in RAM and we measure time

Proposals

- Sadakane's CST `cst_sada`
- Fully Compressed Suffix Tree (Russo et al.)
- CSTs based on interval representation of nodes (Fischer et al. `cstY`, Ohlebusch et al. `cst_sct3`)

CSTs

Goal of a CST implementation

Replace fastest uncompressed ST implementations in different scenarios

- (a) both fit in RAM and we measure time
- (b) both fit in RAM and we measure resource costs
- (c) only CST fits in RAM and we measure time

Proposals **which might work for (a) and (b)**

- **Sadakane's CST** `cst_sada`
- Fully Compressed Suffix Tree (Russo et al.)
- **CSTs based on interval representation of nodes** (Fischer et al. `cstY`, Ohlebusch et al. `cst_sct3`)

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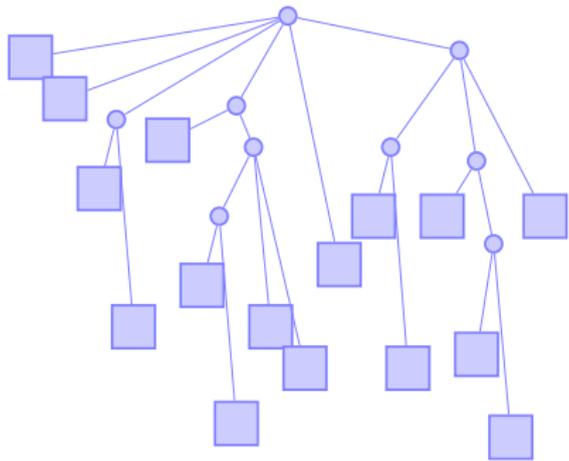
3 CST in practice

- The *sds/* library

Big picture of CST design



Example: Compressing NAV

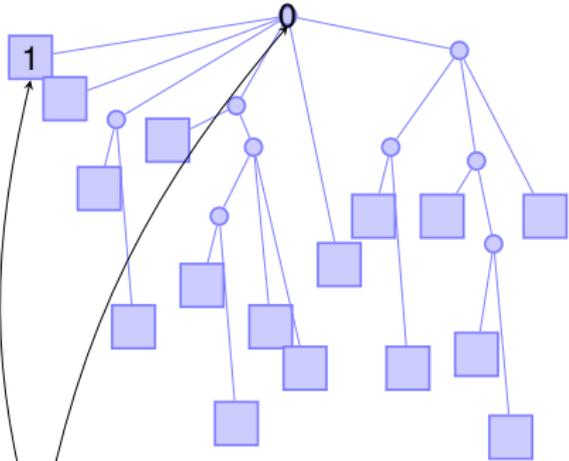


tree uncompressed
 $\mathcal{O}(n \log n)$ bits

$BPS_{dfs} = (00(00)(0((00)00))0((00)(0(00)))0)$

compressed
 $4n$ bits

Example: Compressing NAV

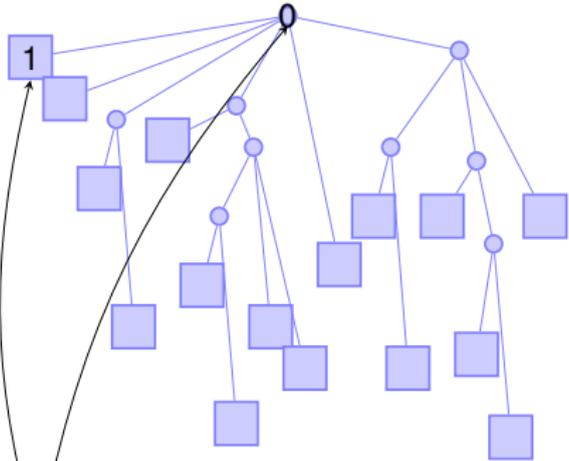


$BPS_{dfs} = ((()())(0((()())0))0((())(0((()()))))$

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Example: Compressing NAV

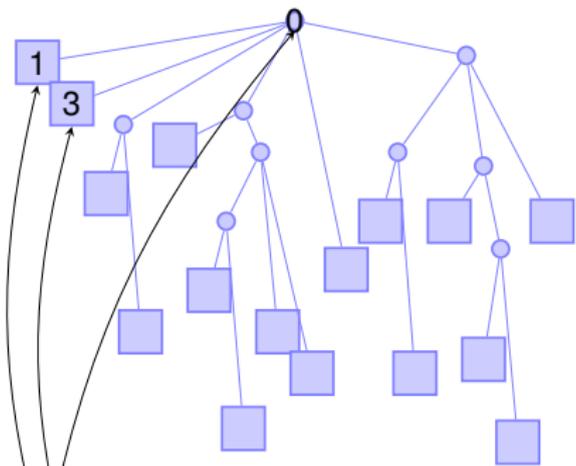


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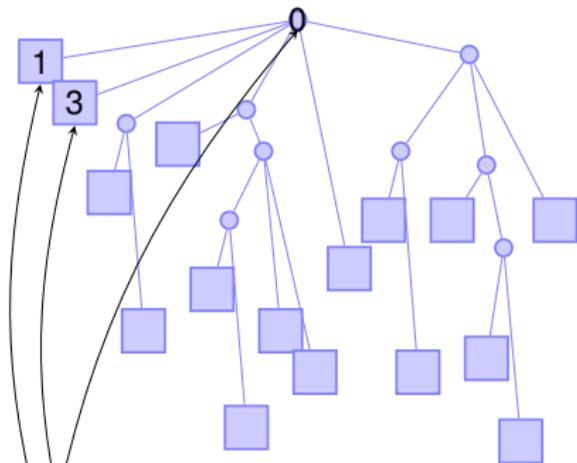


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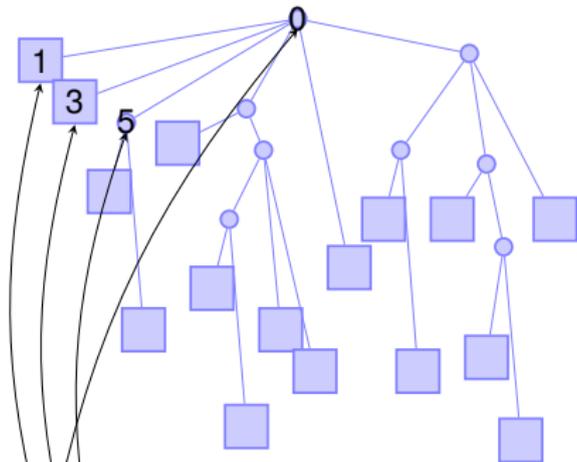


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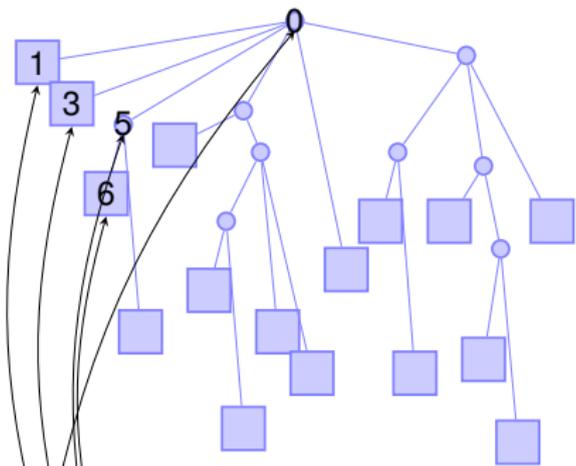


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Example: Compressing NAV



$BPS_{dfs} = (())(())(0((0)0))0((0)(0(0)))$

tree uncompressed
 $O(n \log n)$ bits

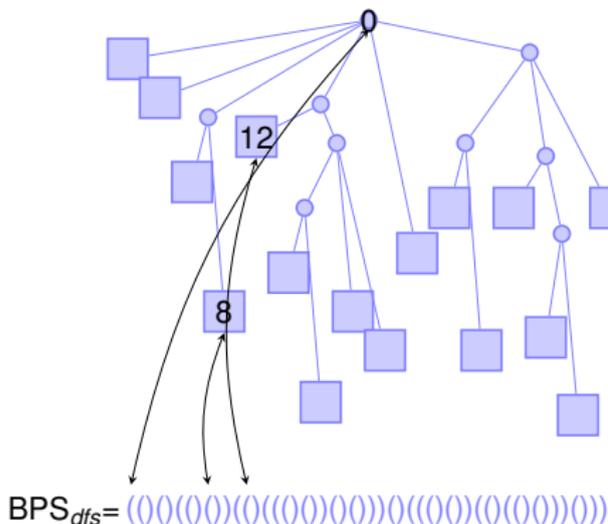
compressed
 $4n$ bits

NAV data structures (2)

Comparison of different NAV structures

space in bits	cst_sada $4n + o(n)$	cst_sct $2n + o(n)$	cst_sct3 $3n + o(n)$
<i>root()</i>	$\mathcal{O}(1)$	$\mathcal{O}(1)$	$\mathcal{O}(1)$
<i>degree(v)</i>	$\mathcal{O}(\sigma)$	$\mathcal{O}(t_{LCP} \log \sigma)$	$\mathcal{O}(1)$
<i>depth(v)</i>	$\mathcal{O}(t_{LCP})$	$\mathcal{O}(t_{LCP})$	$\mathcal{O}(t_{LCP})$
<i>parent(v)</i>	$\mathcal{O}(1)$	$\mathcal{O}(t_{LCP} \log \sigma)$	$\mathcal{O}(1)$
<i>select_child(v, i)</i>	$\mathcal{O}(i)$	$\mathcal{O}(t_{LCP})$	$\mathcal{O}(1)$
<i>sibling(v)</i>	$\mathcal{O}(1)$	$\mathcal{O}(t_{LCP})$	$\mathcal{O}(1)$
<i>sl(v), lca(v, w)</i>	$\mathcal{O}(1)$	$\mathcal{O}(t_{LCP} \log \sigma)$	$\mathcal{O}(1)$
<i>child(v, c)</i>	$\mathcal{O}(t_{SA} \sigma)$	$\mathcal{O}(t_{SA} \log \sigma)$	$\mathcal{O}(t_{SA} \log \sigma)$

Example operations: $select_leaf(i)$ and $lca(v, w)$ on NAV



$$select_leaf(4) = select(4, '10')$$

$$= 8$$

$$select_leaf(5) = select(5, '10')$$

$$= 12$$

$$lca(8, 12) = double_enclose(8, 12)$$

$$= 0$$

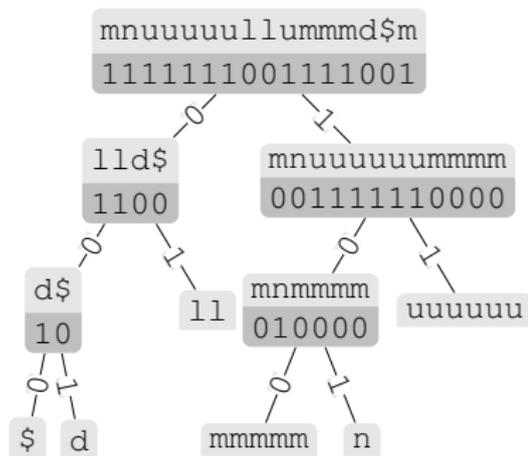
Virtues of a CSA based on BWT

- Small size: $|\text{CSA}| = |\text{BWT}| + \frac{n \log n}{s_{\text{SA}}}$ bits
 where $|\text{BWT}|$ can be chosen to be
 - $n \log \sigma$ bits
 - $nH_0(T)$ bits
 - $nH_k(T) + \mathcal{O}(\sigma^k)$ bits
- pattern matching in time $\mathcal{O}(|P| \log \sigma)$ (even $\mathcal{O}(|P|)$ for $\sigma \in \text{polylog}(n)$) by backward search (Ferragina & Manzini)

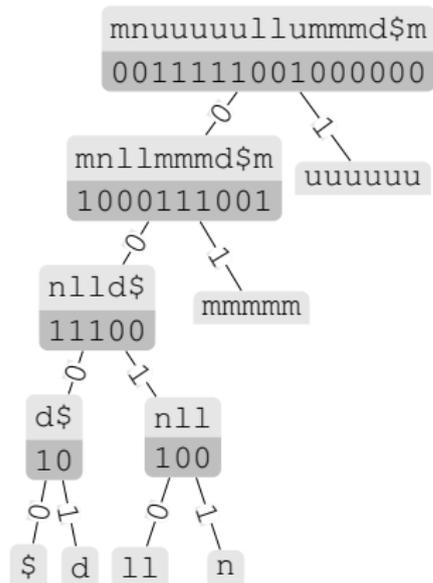
H_k of the *Pizza&Chili* 200MB test cases

k	dblp.xml		dna		english		proteins		rand_k128		sources	
	H_k	CT/n	H_k	CT/n	H_k	CT/n	H_k	CT/n	H_k	CT/n	H_k	CT/n
0	5.257	0.0000	1.974	0.0000	4.525	0.0000	4.201	0.0000	7.000	0.0000	5.465	0.0000
1	3.479	0.0000	1.930	0.0000	3.620	0.0000	4.178	0.0000	7.000	0.0000	4.077	0.0000
2	2.170	0.0000	1.920	0.0000	2.948	0.0001	4.156	0.0000	6.993	0.0001	3.102	0.0000
3	1.434	0.0007	1.916	0.0000	2.422	0.0005	4.066	0.0001	5.979	0.0100	2.337	0.0012
4	1.045	0.0043	1.910	0.0000	2.063	0.0028	3.826	0.0011	0.666	0.6939	1.852	0.0082
5	0.817	0.0130	1.901	0.0000	1.839	0.0103	3.162	0.0173	0.006	0.9969	1.518	0.0250
6	0.705	0.0265	1.884	0.0001	1.672	0.0265	1.502	0.1742	0.000	1.0000	1.259	0.0509
7	0.634	0.0427	1.862	0.0001	1.510	0.0553	0.340	0.4506	0.000	1.0000	1.045	0.0850
8	0.574	0.0598	1.834	0.0004	1.336	0.0991	0.109	0.5383	0.000	1.0000	0.867	0.1255
9	0.537	0.0773	1.802	0.0013	1.151	0.1580	0.074	0.5588	0.000	1.0000	0.721	0.1701
10	0.508	0.0955	1.760	0.0051	0.963	0.2292	0.061	0.5699	0.000	1.0000	0.602	0.2163

Wavelet tree: rank for character sequences



(a)



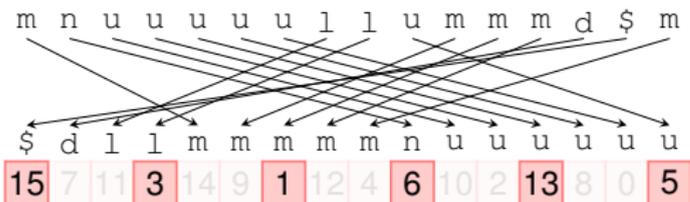
(b)

Example: Compressing CSA (practical approach)

i	SA	LF	T^{BWT}	T
0	15	4	m	\$
1	7	9	n	dumulumum\$
2	11	10	u	lmum\$
3	3	11	u	lmundumulumum\$
4	14	12	u	m\$
5	9	13	u	mulmum\$
6	1	14	u	mulmundumulumum\$
7	12	2	l	mum\$
8	4	3	l	mundumulumum\$
9	6	15	u	ndumulumum\$
10	10	5	m	ulumum\$
11	2	6	m	ulmundumulumum\$
12	13	7	m	um\$
13	8	1	d	umulumum\$
14	0	0	\$	umulumundumulumum\$
15	5	8	m	undumulumum\$

Example: Compressing CSA (practical approach)

0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
15	7	11	3	14	9	1	12	4	6	10	2	13	8	0	5



$$s_{SA} = 3$$

access LF[i] in time $\mathcal{O}(\log \sigma)$

access CSA[i] in time $\mathcal{O}(s_{SA} \log \sigma)$

SA uncompressed
 $n \log n$ bits

compressed

CSA = T^{BWT} + SA samples:

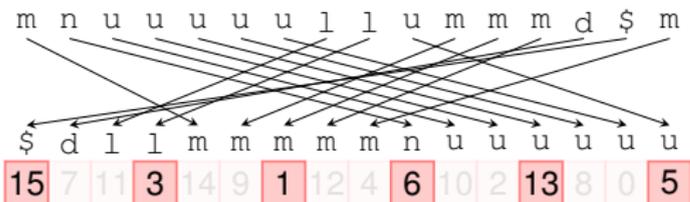
$n \log \sigma + o(n \log \sigma)$ bits

+
 $\frac{n \log n}{s_{SA}}$ bits

Example: Compressing CSA (practical approach)

0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
15	7	11	3	14	9	1	12	4	6	10	2	13	8	0	5

SA[13]=



$s_{SA} = 3$

access LF[i] in time $\mathcal{O}(\log \sigma)$

access CSA[i] in time $\mathcal{O}(s_{SA} \log \sigma)$

SA uncompressed
 $n \log n$ bits

compressed

CSA = T^{BWT} + SA samples:

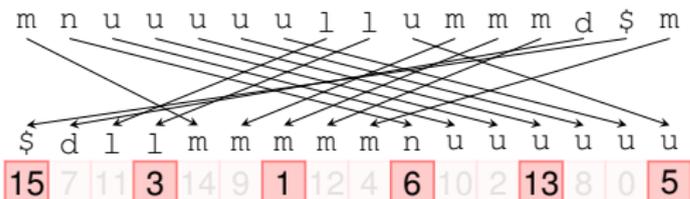
$n \log \sigma + o(n \log \sigma)$ bits

+
 $\frac{n \log n}{s_{SA}}$ bits

Example: Compressing CSA (practical approach)

0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
15	7	11	3	14	9	1	12	4	6	10	2	13	8	0	5

$$SA[13] = SA[1] + 1$$



$$s_{SA} = 3$$

access LF[i] in time $\mathcal{O}(\log \sigma)$

access CSA[i] in time $\mathcal{O}(s_{SA} \log \sigma)$

SA uncompressed
 $n \log n$ bits

compressed

CSA = T^{BWT} + SA samples:

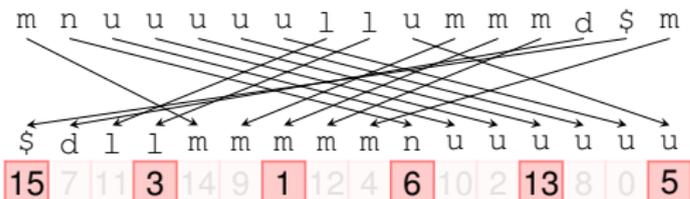
$n \log \sigma + o(n \log \sigma)$ bits

+
 $\frac{n \log n}{s_{SA}}$ bits

Example: Compressing CSA (practical approach)

0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
15	7	11	3	14	9	1	12	4	6	10	2	13	8	0	5

$$SA[13]=6+2=8$$



$$s_{SA} = 3$$

access LF[i] in time $\mathcal{O}(\log \sigma)$

access CSA[i] in time $\mathcal{O}(s_{SA} \log \sigma)$

SA uncompressed
 $n \log n$ bits

compressed

CSA = T^{BWT} + SA samples:

$n \log \sigma + o(n \log \sigma)$ bits

+
 $\frac{n \log n}{s_{SA}}$ bits

Overview of LCP data structures

data structure	uses	access	memory in bits
<code>lcp_uncompressed</code>	-	$\mathcal{O}(1)$	$n \log n$
<code>lcp_support_sada</code>	CSA	$\mathcal{O}(t_{\text{SA}})$	$2n + o(n)$
<code>lcp_kurtz</code>	-	$\mathcal{O}(\log n)$ or $\mathcal{O}(1)$	$8n_1 + 2n_2 \log n$
...
<code>lcp_support_tree</code>	NAV	$\mathcal{O}(\log \sigma')$	$H_0 q_1 + q_2 \log n$
<code>lcp_support_tree2</code>	NAV & LF	$\mathcal{O}(s_{\text{LCP}} \log \sigma')$	$H_0 q_1 + (q_2 \log n) / s_{\text{LCP}}$

with $n_1 + n_2 = n$

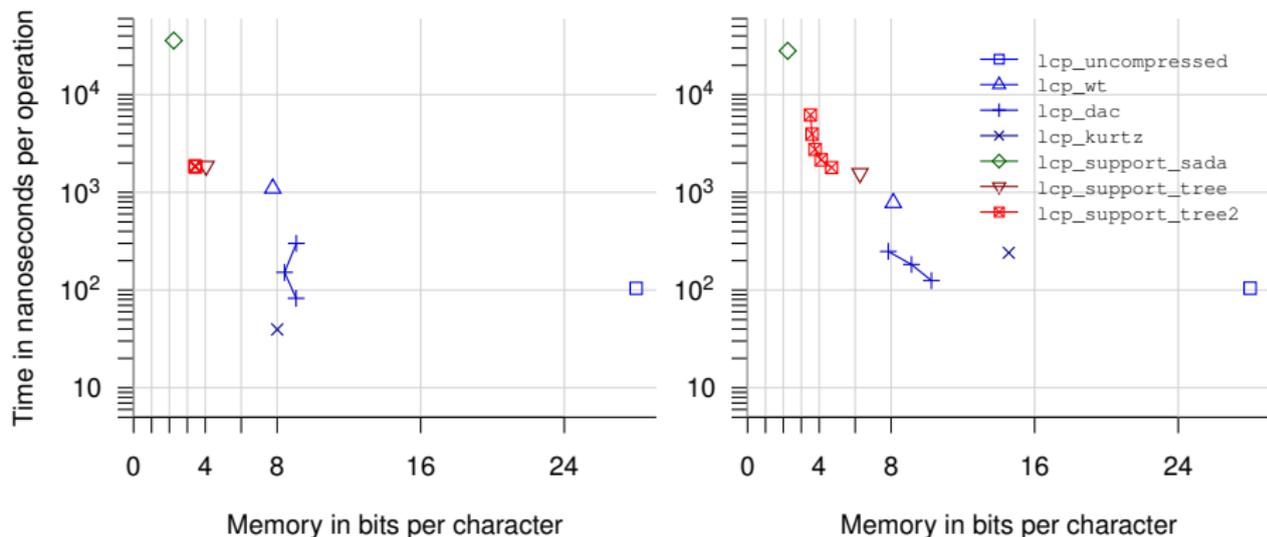
and $q_1 + q_2 = q < n$

q number of inner nodes of the ST

Runtime for random access to the LCP array (1)

dblp.xml.200MB

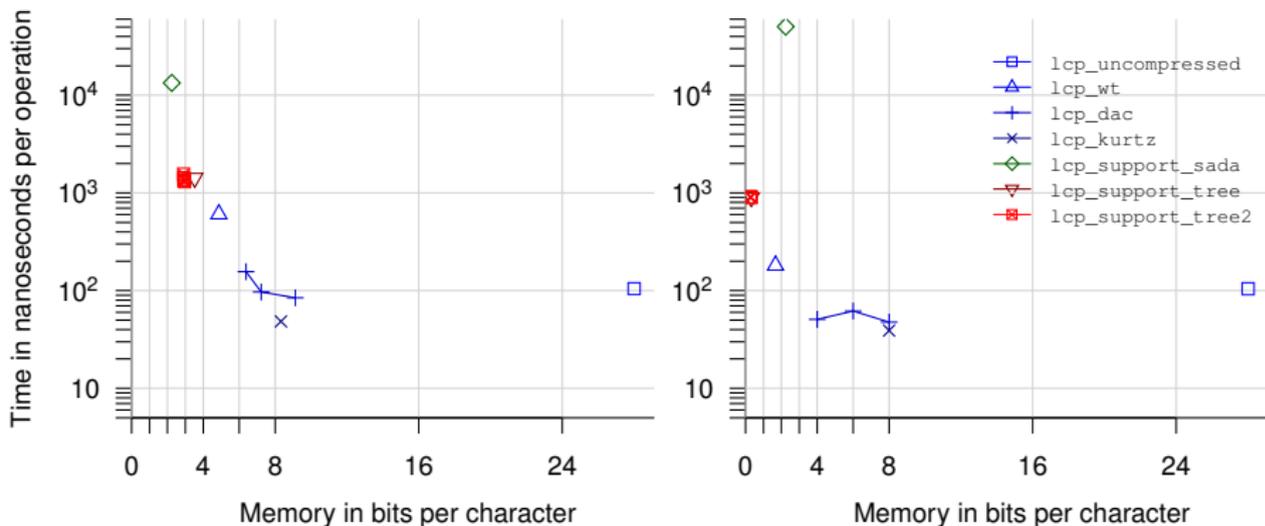
proteins.200MB



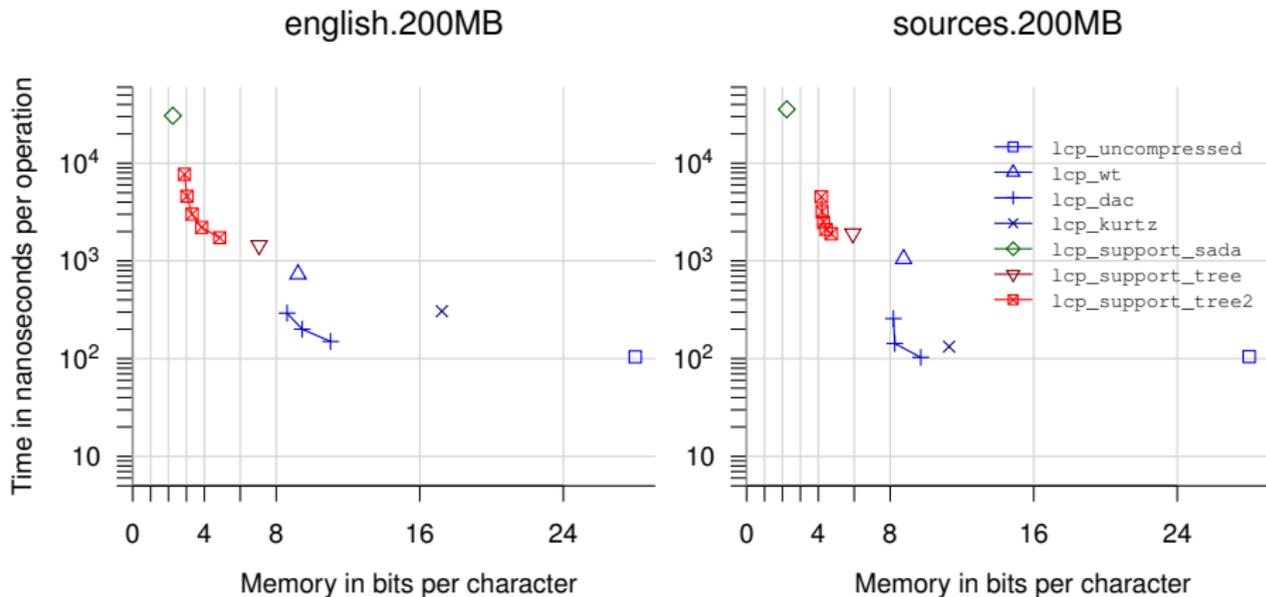
Runtime for random access to the LCP array (2)

dna.200MB

rand_k128.200MB



Runtime for random access to the LCP array (3)



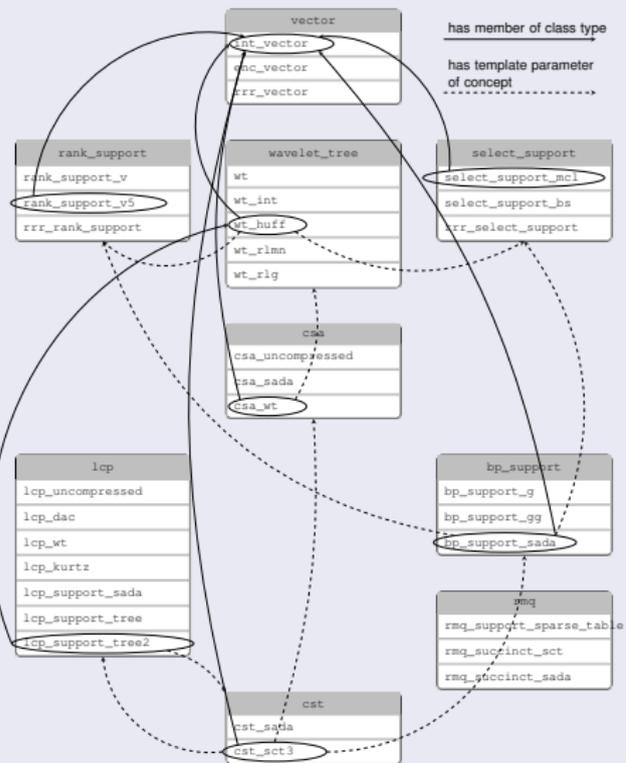
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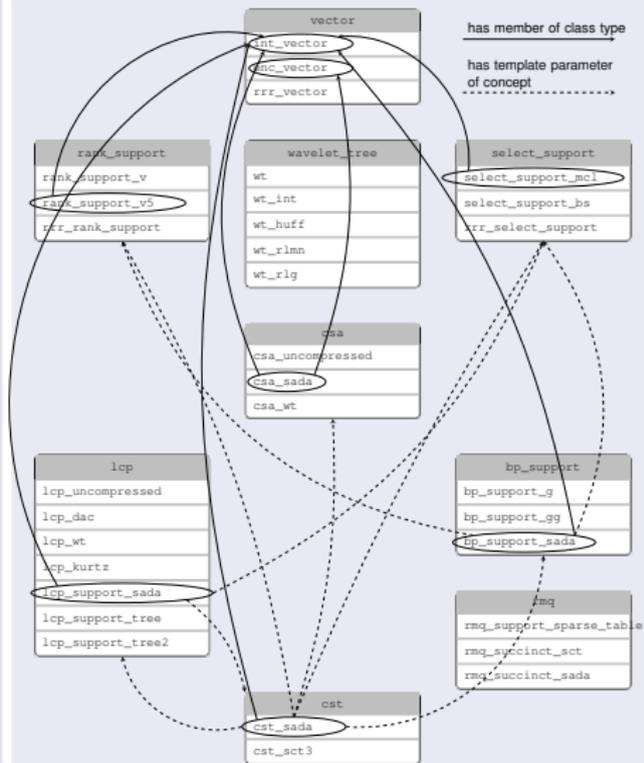
The succinct data structure library *sds/*

- Provides basic and advanced succinct data structures
- Easy to use (very similar to C++ STL)
- Fast and space-efficient construction of data structures
- 64-bit implementation
- Well-optimized implementation (e.g. now using hardware POPCOUNT operation,..)
- Easy configuration of myriads of CSAs, CSTs with many time-space trade-offs
- Fast prototyping of other complex succinct data structures

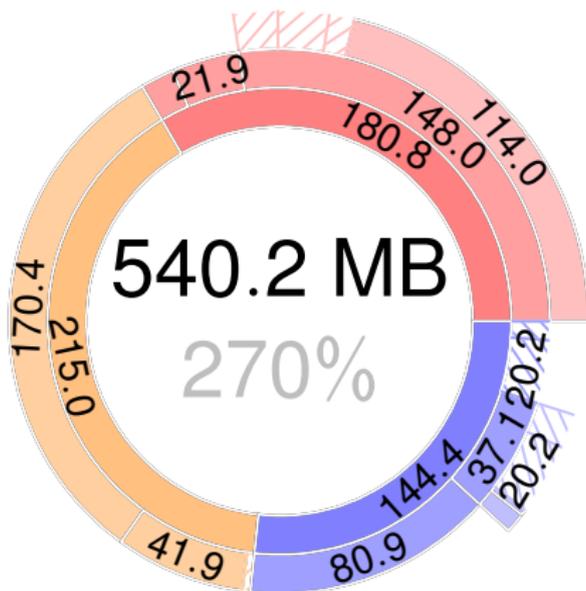
cst_sada<csa_sada<>, lcp_..



cst_sct3<csa_wt<wt_huff<>..



CST space in practice (for english.200MB)



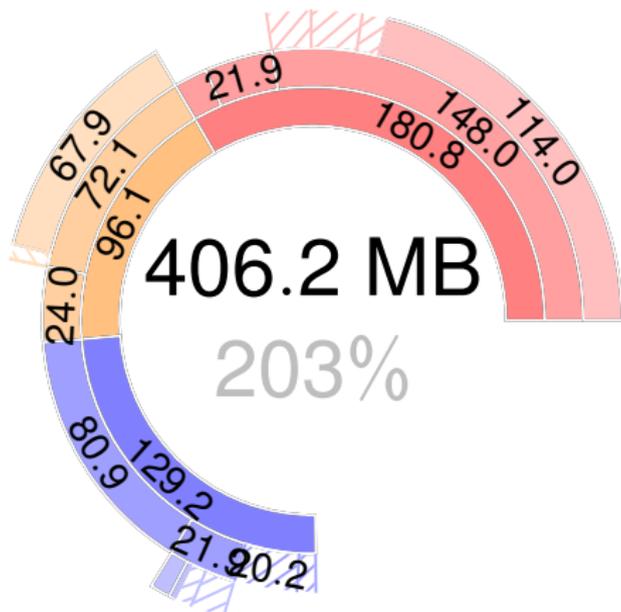
```

CSA: 180.8 MB
wt: 148 MB
  data: 114 MB
  rank: 7.1 MB
  select 1: 14.4 MB
  select 0: 12.5 MB
sa_sample: 21.9 MB
isa_sample: 10.9 MB
lcp: 215 MB
  lcp values: 170.4 MB
  overflow mark: 41.9 MB
rank: 2.6 MB
nav: 144.4 MB
BPSoffs(bit_vector): 80.9 MB
bp_support: 37.1 MB
  small block: 5.7 MB
  medium block: 1.6 MB
bp rank: 20.2 MB
bp select: 9.6 MB
rank_support10: 20.2 MB
select_support10: 6.1 MB

```

```
cst_sada<csa_wt<wt_huff<> >,lcp_dac<> >
```

CST space in practice (for english.200MB)

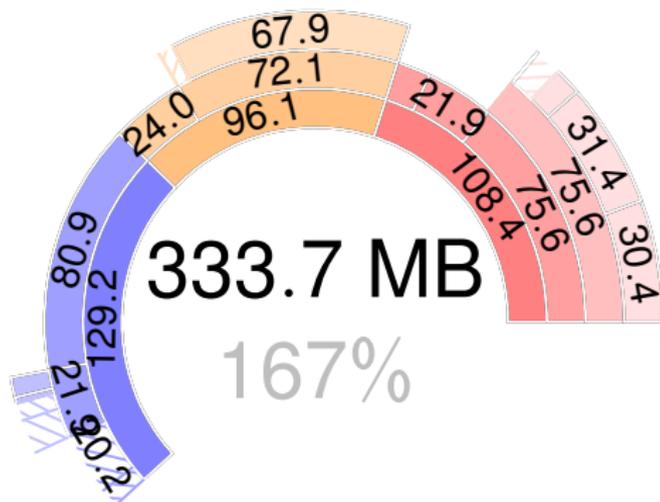


```

CSA: 180.8 MB
wt: 148 MB
  data: 114 MB
  rank: 7.1 MB
  select 1: 14.4 MB
  select 0: 12.5 MB
sa_sample: 21.9 MB
isa_sample: 10.9 MB
lcp: 96.1 MB
  small lcp: 72.1 MB
  data: 67.9 MB
  rank: 4.2 MB
  big lcp: 24 MB
nav: 129.2 MB
  bp: 80.9 MB
  bp_support: 21.9 MB
  small block: 5.7 MB
  medium block: 1.6 MB
  bp rank: 5.1 MB
  bp select: 9.6 MB
rank_support10: 20.2 MB
select_support10: 6.1 MB
  
```

```
cst_sada<csa_wt<wt_huff<> >,lcp_support_tree2<> >
```

CST space in practice (for english.200MB)



```

CSA: 108.4 MB
wt: 75.6 MB
data: 75.6 MB
bt: 30.4 MB
btnr: 31.4 MB
btnrp: 6.7 MB
rank samples: 6.9 MB
invert: 0.2 MB
rank: 0 MB
select 1: 0 MB
select 0: 0 MB
sa_sample: 21.9 MB
isa_sample: 10.9 MB
lcp: 96.1 MB
small lcp: 72.1 MB
data: 67.9 MB
rank: 4.2 MB
big lcp: 24 MB
nav: 129.2 MB
BPSdfs(bit_vector): 80.9 MB
bp_support: 21.9 MB
small block: 5.7 MB
medium block: 1.6 MB
bp rank: 5.1 MB
bp select: 9.6 MB
rank_support10: 20.2 MB
select_support10: 6.1 MB

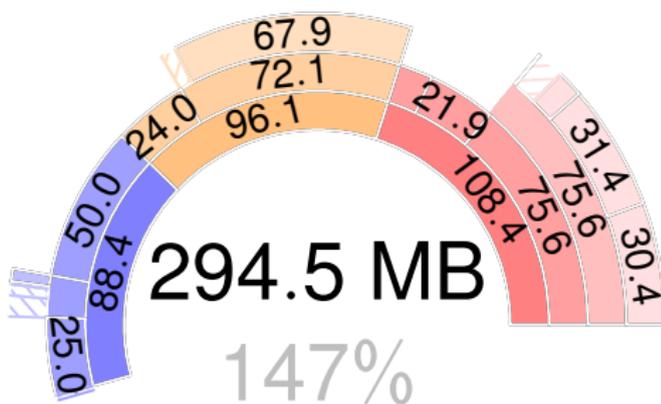
```

```

cst_sada<csa_wt<wt_huff<rrr_vector<> >
>,lcp_support_tree2<> >

```

CST space in practice (for english.200MB)



```

CSA: 108.4 MB
wt: 75.6 MB
  data: 75.6 MB
    bt: 30.4 MB
    btr: 31.4 MB
    btrp: 6.7 MB
    rank samples: 6.9 MB
    invert: 0.2 MB
  rank: 0 MB
  select 1: 0 MB
  select 0: 0 MB
sa_sample: 21.9 MB
isa_sample: 10.9 MB
lcp: 96.1 MB
  small lcp: 72.1 MB
    data: 67.9 MB
    rank: 4.2 MB
  big lcp: 24 MB
nav: 88.4 MB
bp: 50 MB
bp_support: 13.4 MB
  small block: 3.5 MB
  medium block: 0.8 MB

```

```

cst_sct3<csa_wt<wt_huff<rrr_vector<> >
>,lcp_support_tree2<> >

```

Experimental setup

0 $\hat{=}$ cst_sada<csa_sada<>, lcp_dac<> >

1 $\hat{=}$ cst_sada<csa_sada<>, lcp_support_tree2<> >

2 $\hat{=}$ cst_sada<csa_wt<>, lcp_dac<> >

3 $\hat{=}$ cst_sada<csa_wt<>, lcp_support_tree2<> >

4 $\hat{=}$ cst_sct3<csa_sada<>, lcp_dac<> >

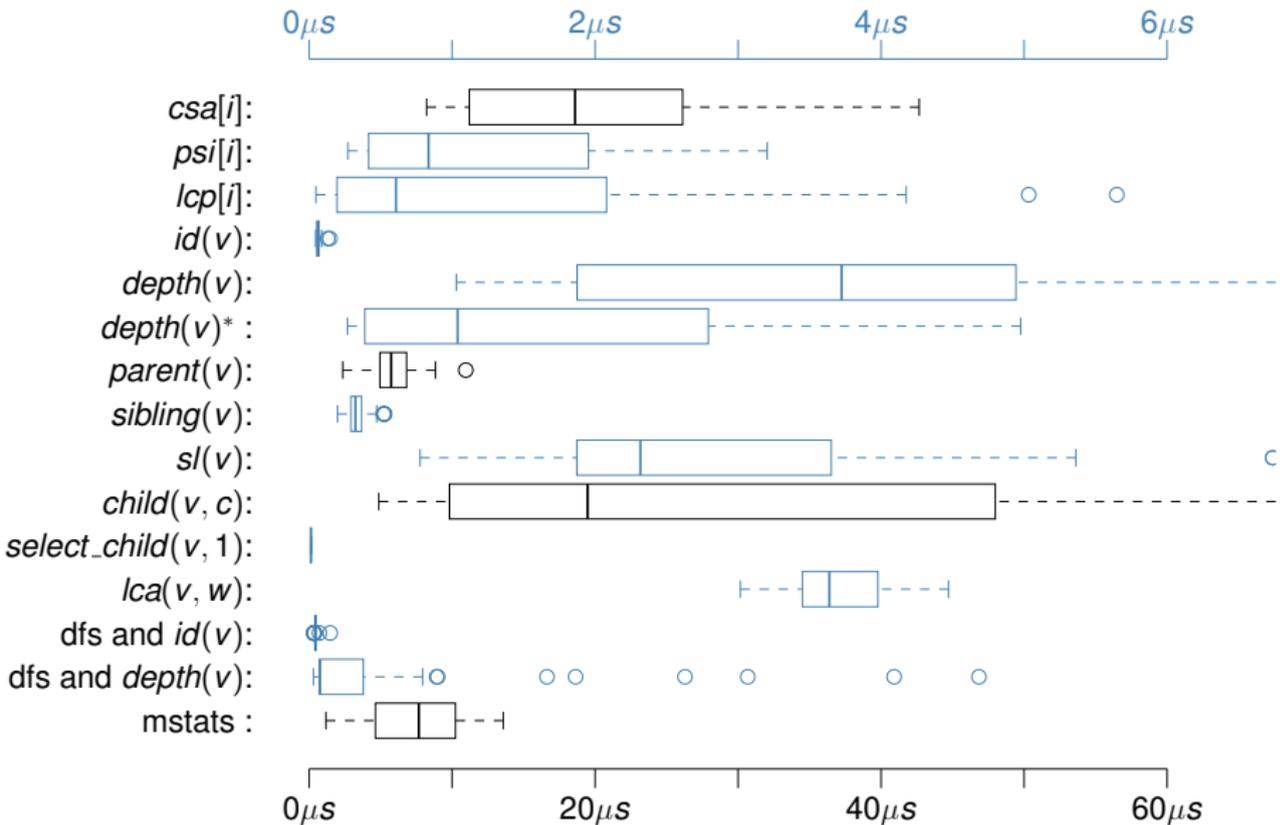
5 $\hat{=}$ cst_sct3<csa_sada<>, lcp_support_tree2<> >

6 $\hat{=}$ cst_sct3<csa_wt<>, lcp_dac<> >

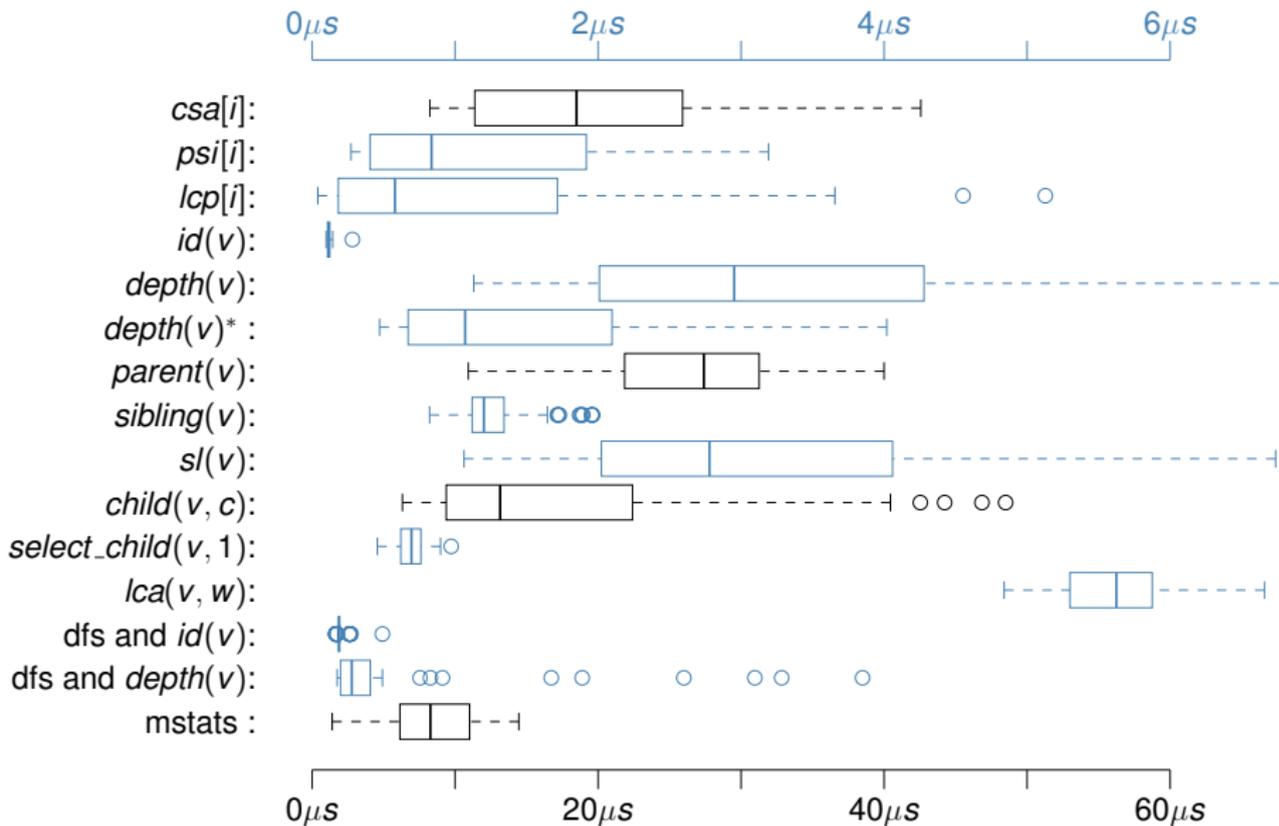
7 $\hat{=}$ cst_sct3<csa_wt<>, lcp_support_tree2<> >

The same basic data structures are used, i.e. its a very fair comparison

Runtime of operations of `cst_sada`

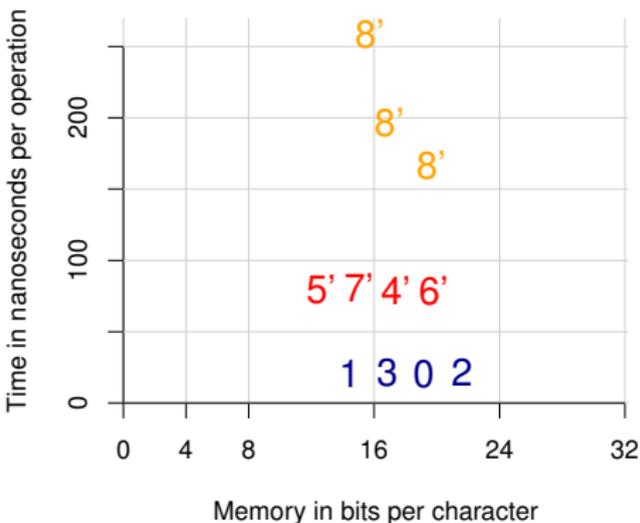


Runtime of operations of `cst_sct3`

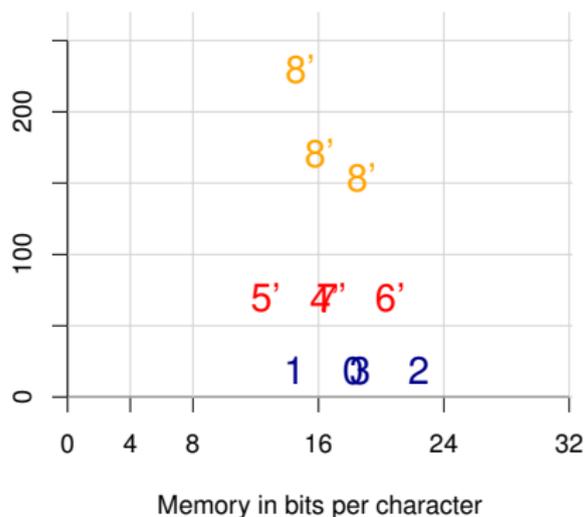


Time-space trade-off for *select_child(v, 1)*

english.200MB

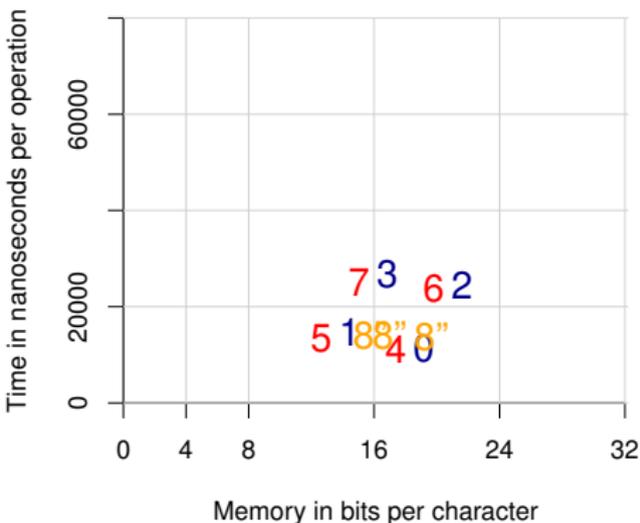


sources.200MB

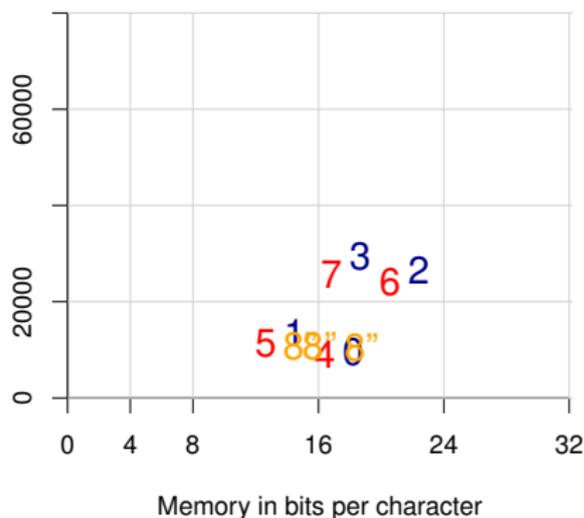


Time-space trade-off for $child(v, c)$

english.200MB



sources.200MB



Construction of a CST

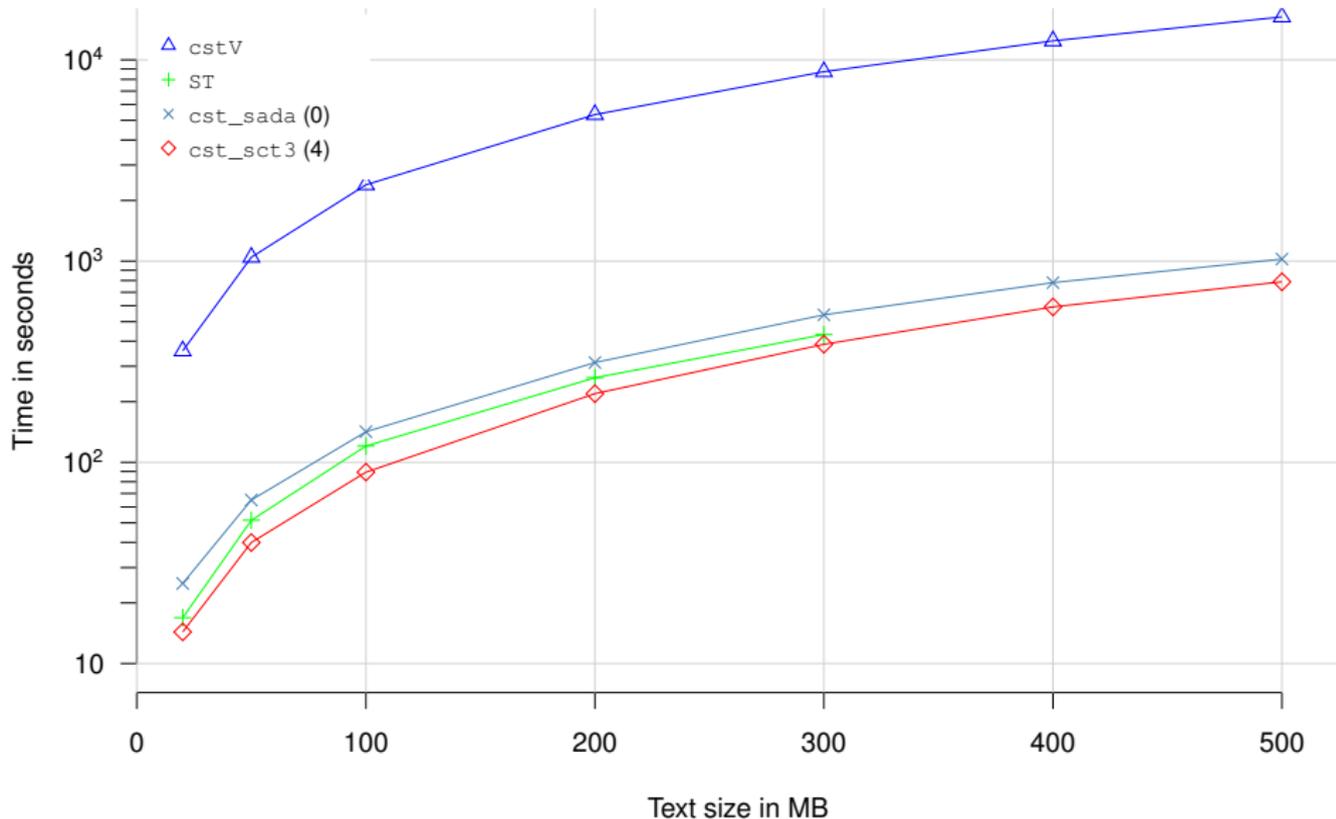
```
#include <sdsl/suffixtrees.hpp>
#include <sdsl/util.hpp>

using namespace sdsl;

typedef cst_sct3<> tCST;

int main(int argc, char* argv[]){
    tCST cst;
    construct_cst( argv[1], cst );
}
```

Runtime for construction (prefixes of english text)

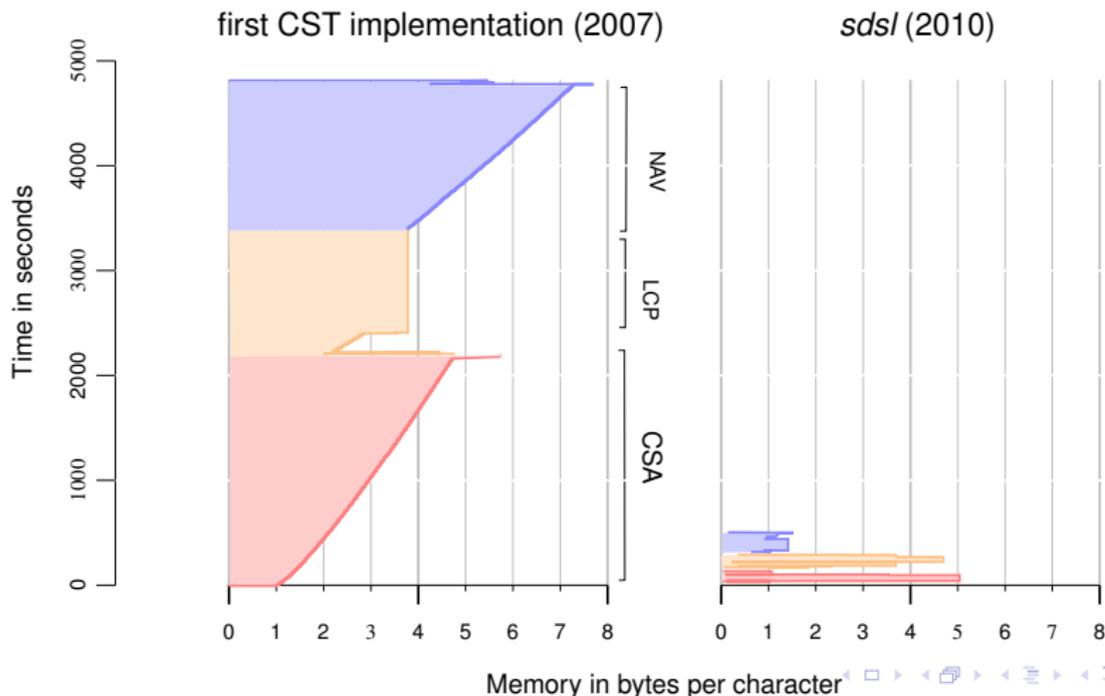


CST construction – resources comparison

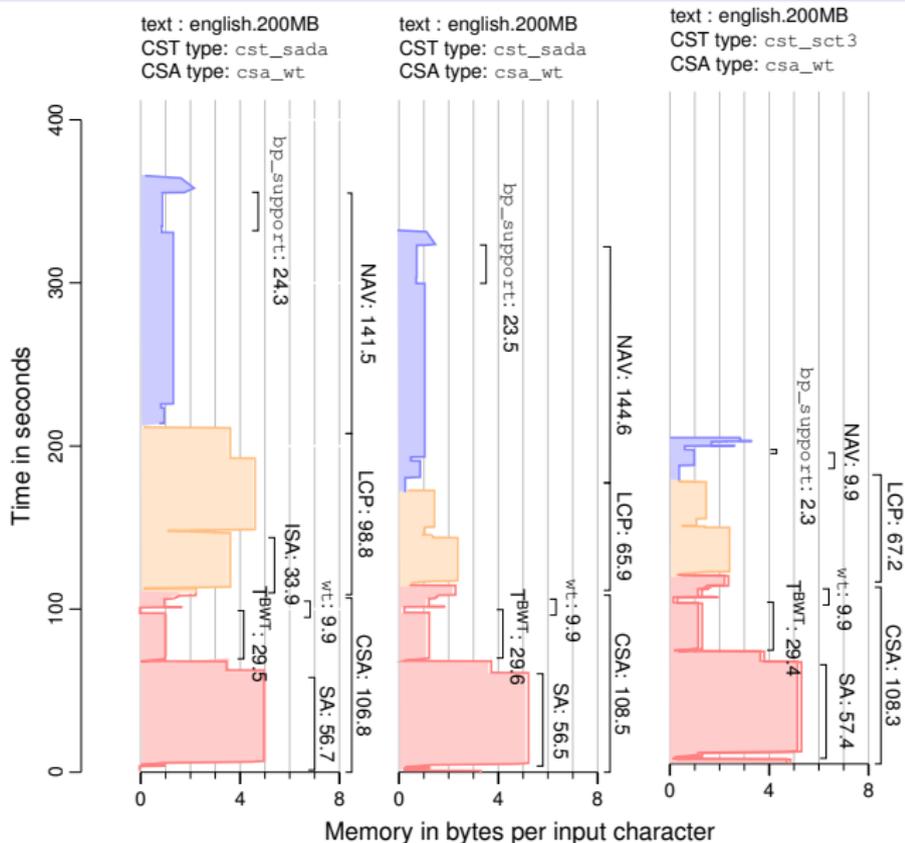
text: english.200MB
 σ : 226

CST type: cst_sada
LCP type: lcp_sada

CSA type: csa_wt



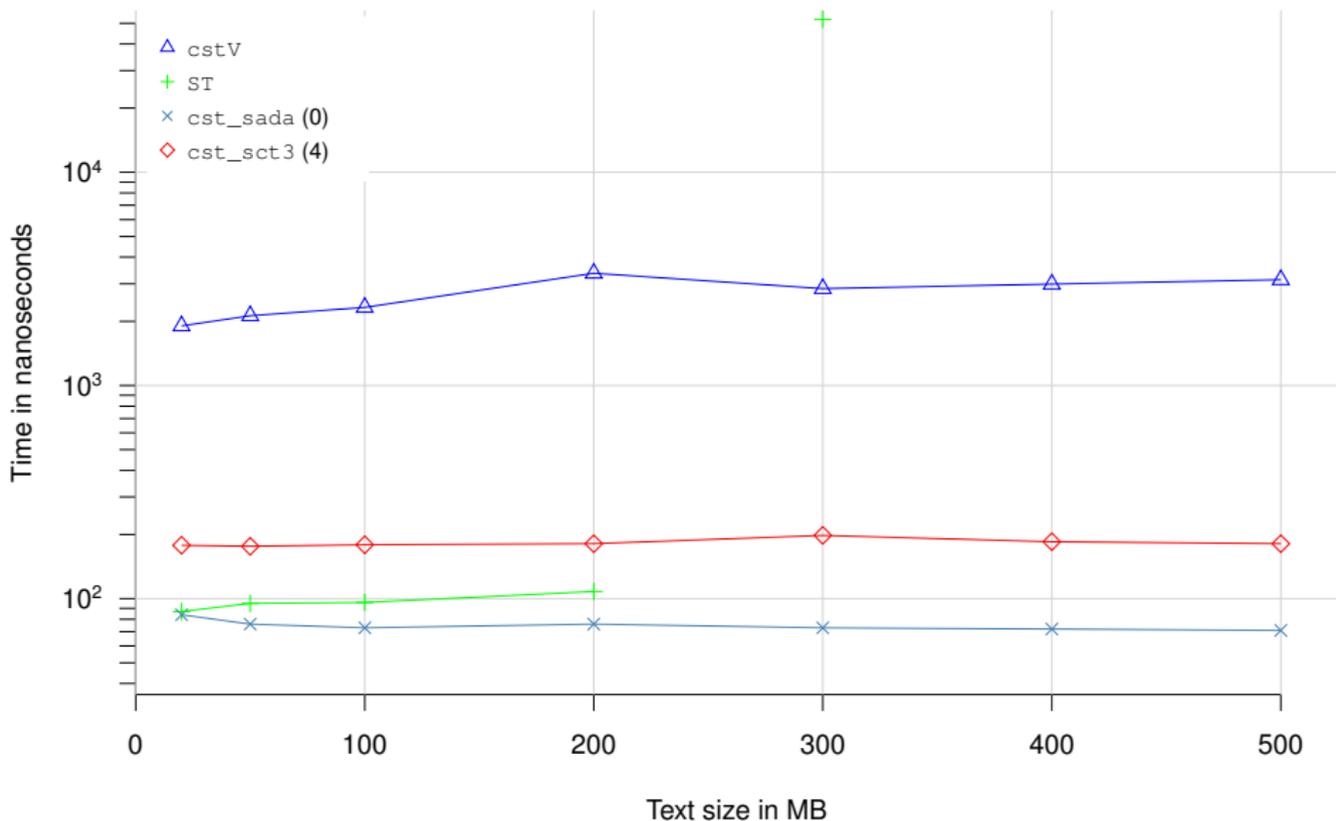
Detailed resources for the construction of CSTs



Depth first search traversal in a CST

```
template<class Cst>
void test_cst_dfs_iterator_and_depth(Cst &cst){
    typedef typename Cst::const_iterator iterator;
    long long cnt = 0;
    for(iterator it=cst.begin(); it!=cst.end();++ it ){
        if( !cst.is_leaf(* it) )
            cnt += cst.depth(* it);
    }
    cout << cnt << endl;
}
```

Runtime of dfs on CST (prefixes of english text)



Conclusion

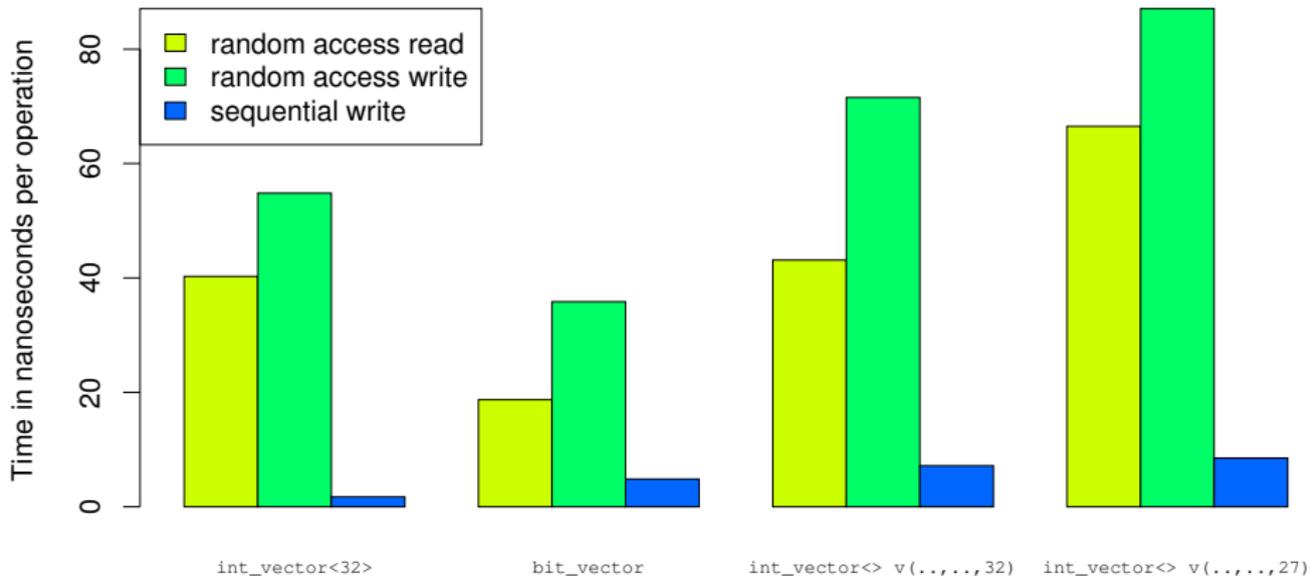
CSTs ...

- ... can be build fast and space-efficient
- ... provide a rich set of functionality
 - fast operations: basic navigation, access LF or Ψ
 - slow operations: $child(v, c)$, access to SA

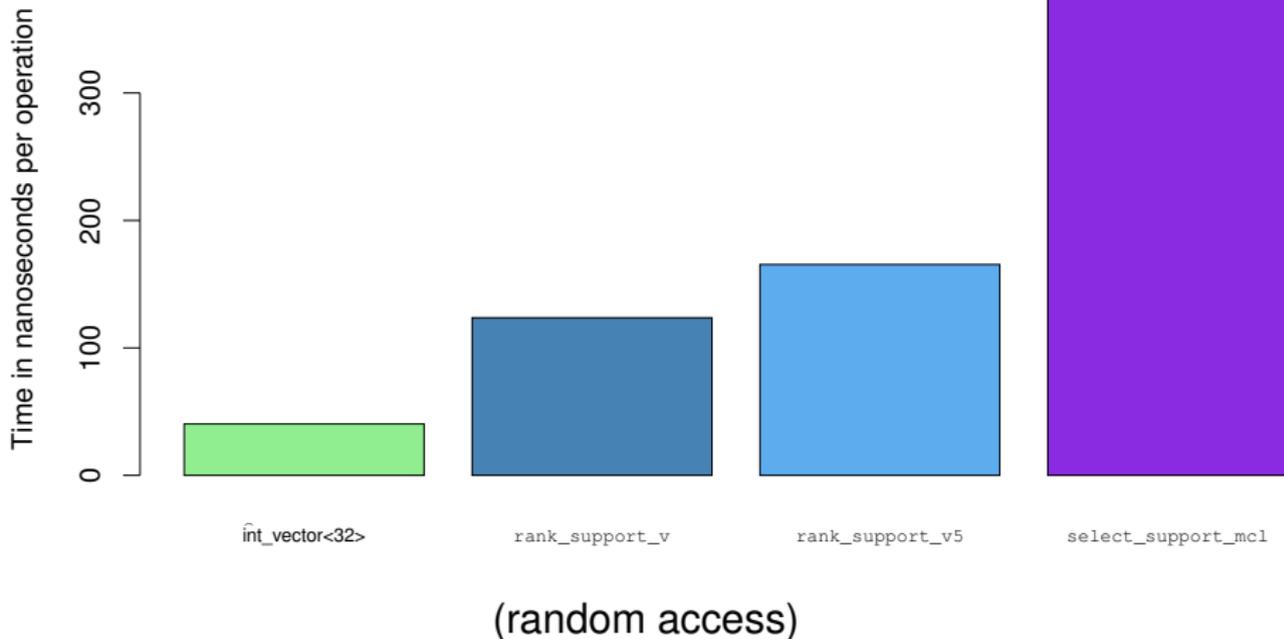
You can use the *sds/* library to configure a CSTs which fits your needs

Thank you!

Runtime of `int_vector` access

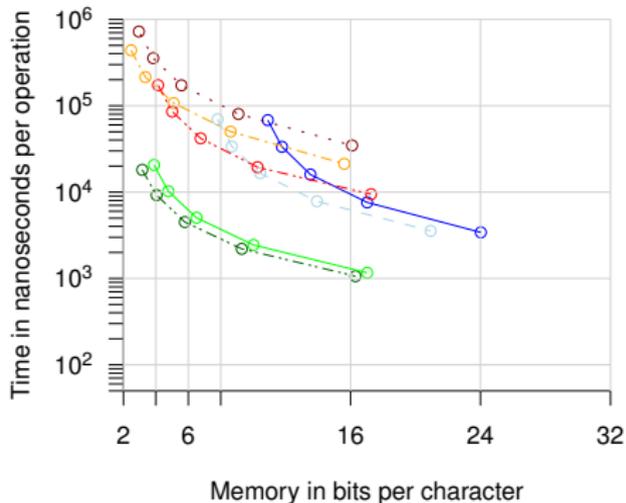


Operation runtime of basic data structures

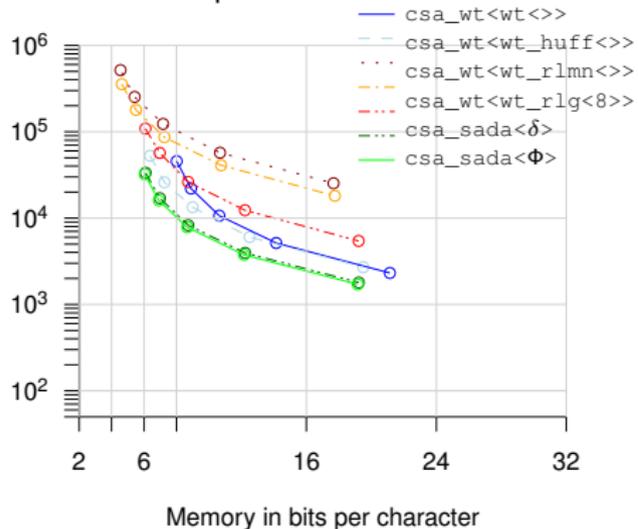


Runtime of CSA access (1)

dblp.xml.200MB

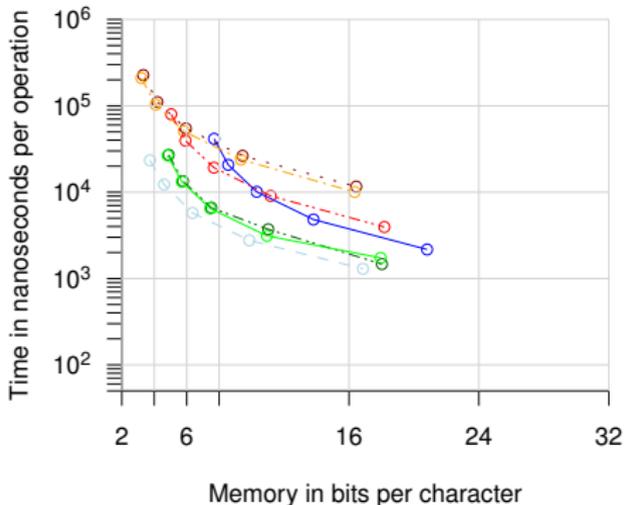


proteins.200MB

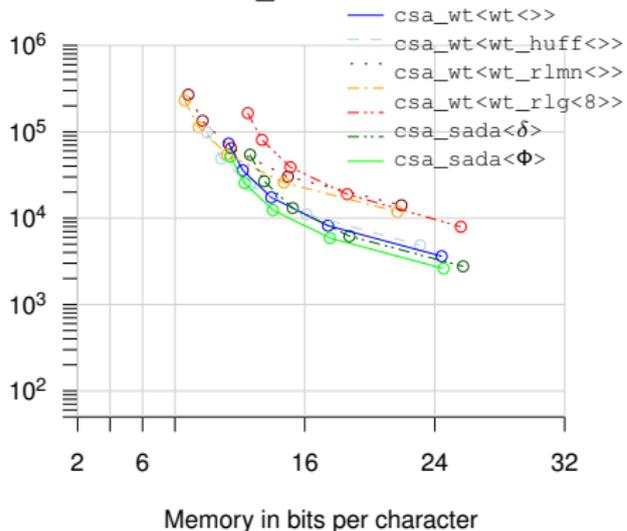


Runtime of CSA access (2)

dna.200MB

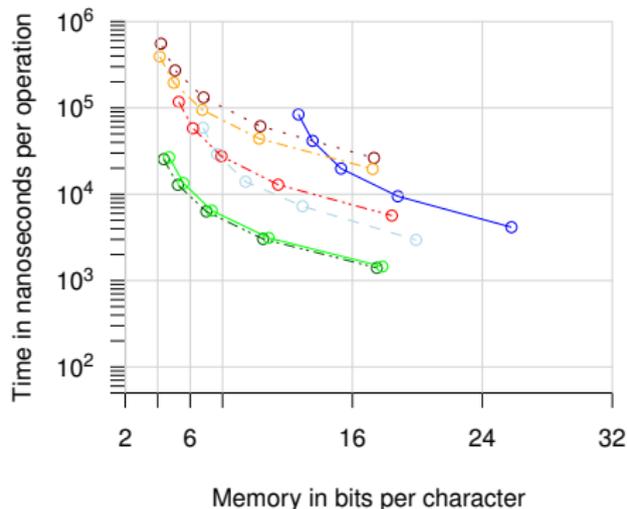


rand_k128.200MB

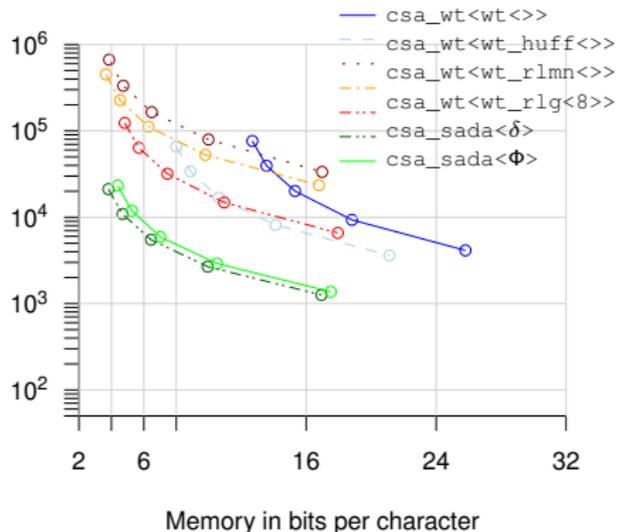


Runtime of CSA access (3)

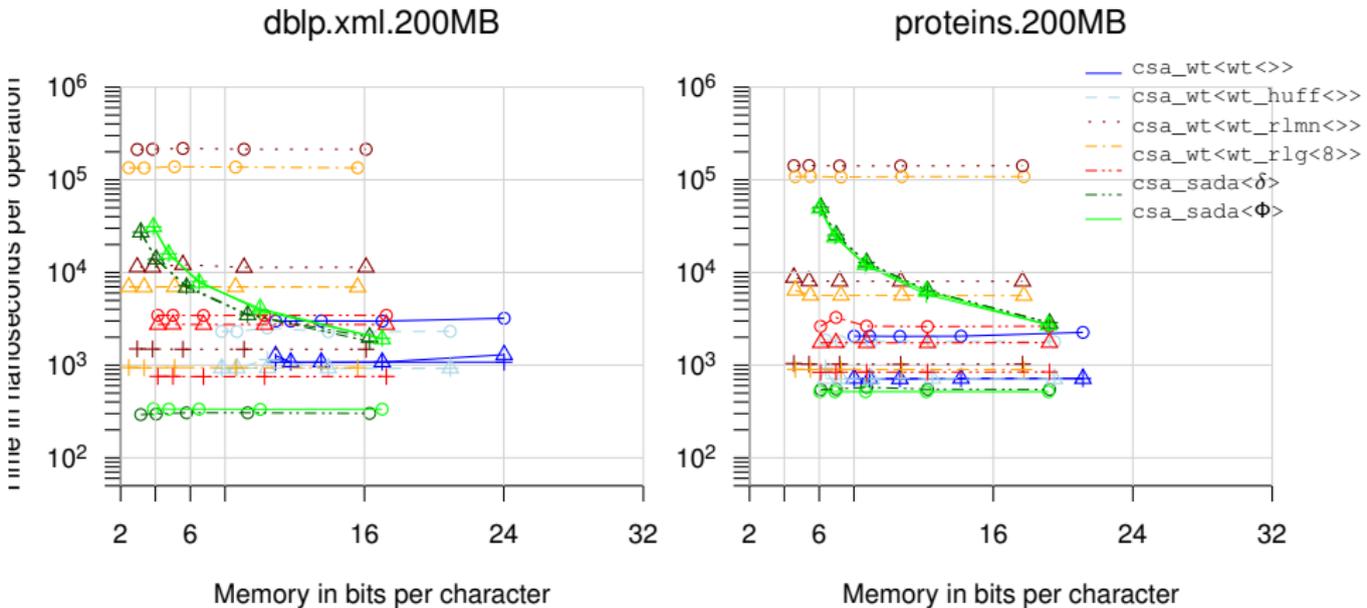
english.200MB



sources.200MB



Runtime of CSA operations (1)

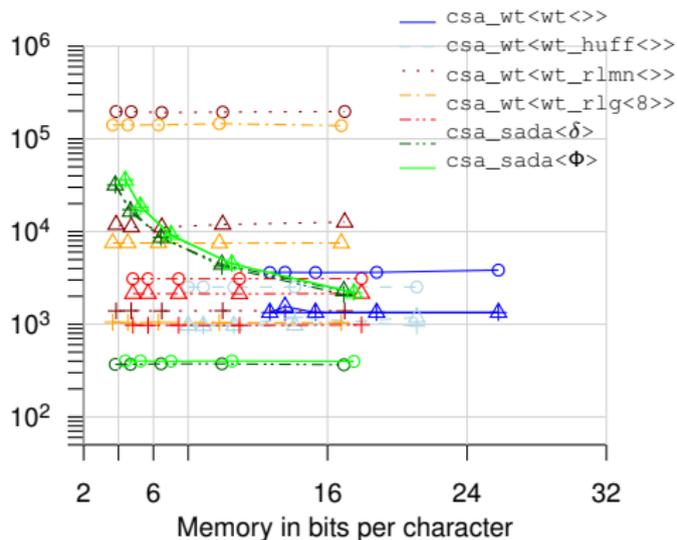
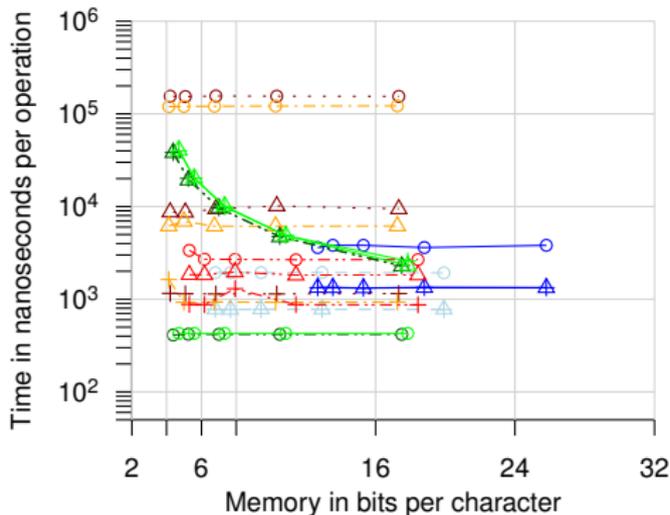


$\circ = \text{psi}[i], \Delta = \text{psi}(i) = \text{LF}[i], + = \text{bwt}[i]$

Runtime of CSA operations (3)

english.200MB

sources.200MB



○ = $\psi[i]$, Δ = $\psi(i) = LF[i]$, + = $bwt[i]$