

Look at that sequence... Is it a vector? Is it a list?
No! It's a Super Tree!!

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indizen  using std::cpp

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Universidad Carlos III de Madrid, November the 30th 2017

Presentation available in my semiabandoned blog:
<http://www.mkrevuelta.com>

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Introduction to the problem

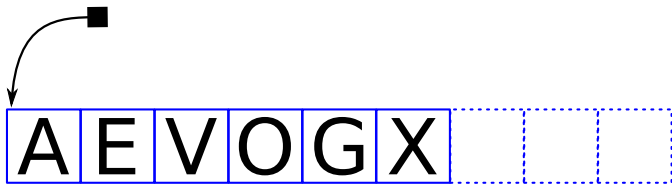
Are lists evil?—Bjarne Stroustrup



<https://isocpp.org/blog/2014/06/stroustrup-lists>

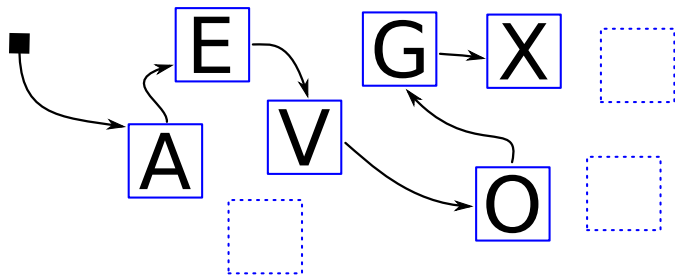
Array

- Random access is fast
- Insertion/extraction are... slow

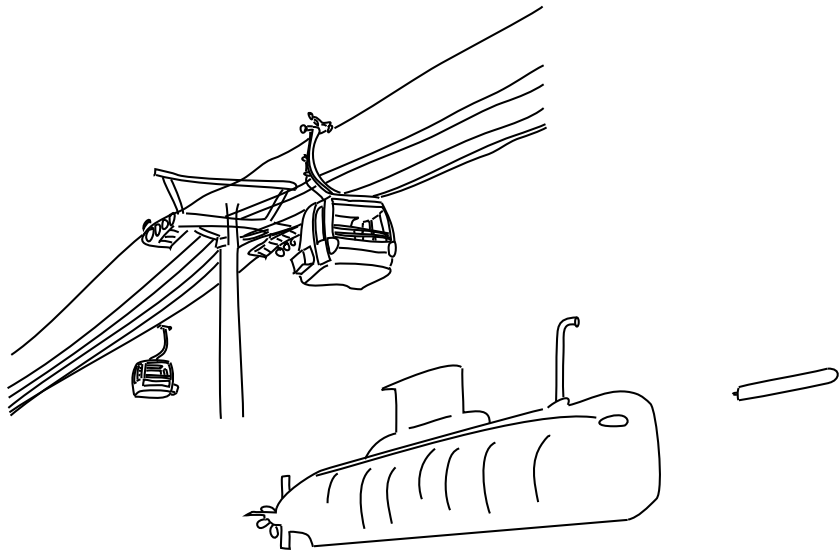


Linked list

- Insertion/extraction are fast
- Random access is... sloooow



How to compare them?



Jon Bentley's suggestion



```
for (;;)
{
```

Random
access

Insertion /
extraction

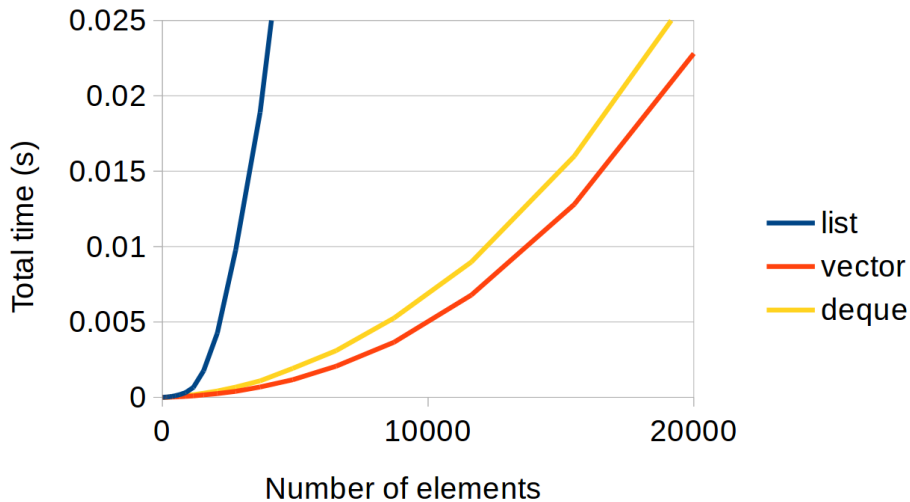
```
}
```

Jon Bentley's suggestion

“Insert a sequence of random integers
into a sorted sequence,

then **remove** those elements **one by one**
as determined by
a **random** sequece of **positions**”

Results



Conclusion

Vectors are faster
by some fixed proportion
(a considerable proportion)

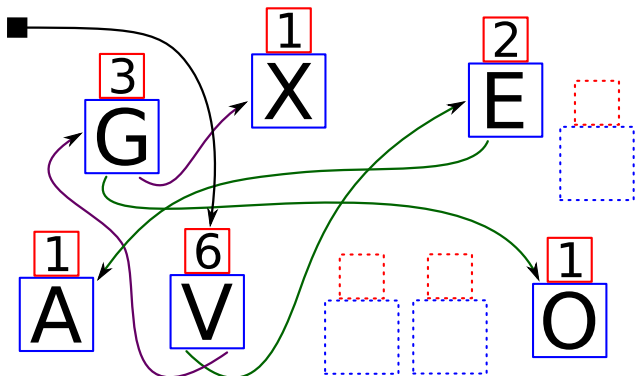
But...

Are we really interested
in Jon Bentley's problem?

Super Tree

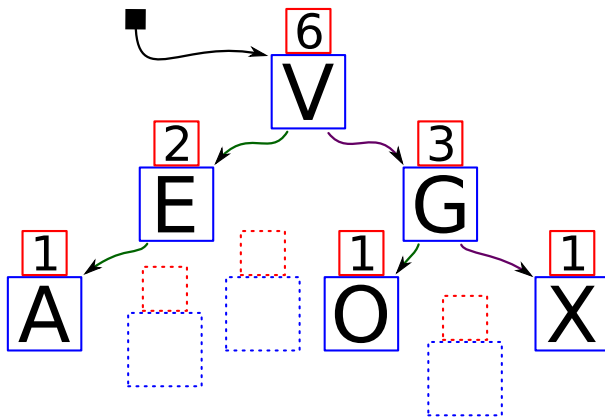
Augmented tree (messed up)

Like a list, but with two “next”s (left, and right)



Augmented tree

Special metadata: number of nodes in the sub-tree



Random access (1/3)

```
template <typename T>
struct node
{
    node<T> * left;        // Left sub-tree
    node<T> * right;      // Right "
    std::size_t count;    // Num. of nodes
    T value;              // Payload
};
```


Random access (2/3)

```
template <typename T>
node<T> * RandomAccess (node<T> * root,
                       std::size_t pos)
{
    if (pos >= root->count)
        return nullptr;

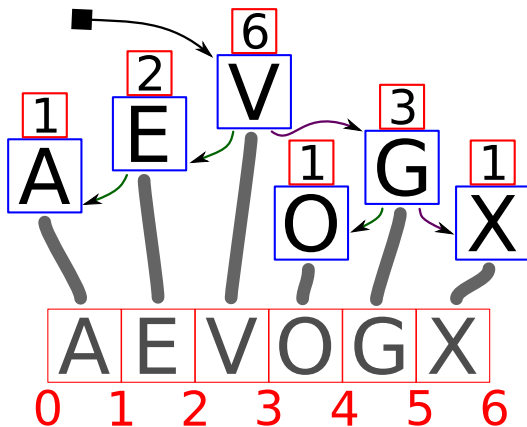
    node<T> * p = root;
```

Random access (3/3)

```
for (;;)
{
    std::size_t nLeft = p->left ?
                        p->left->count : 0;

    if (pos == nLeft)        return p;
    else if (pos < nLeft)   p = p->left;
    else // (pos > nLeft)
    {
        pos -= nLeft + 1;
        p = p->right;
    }                       } } // end
```

Proportional view



Computational complexity










	Random access	Insertion/ Extraction	Sum of both
Array	$O(1)$	$O(N)$	$O(N)$
List	$O(N)$	$O(1)$	$O(N)$
Super Tree	$O(\log(N))$	$O(\log(N))$	$O(\log(N))$

Computational complexity (legend)







- $O(1)$ = constant 🤖
- $O(\log(N))$ = logarithmic 😊
- $O(N)$ = linear 😐
- $O(N \log(N))$ = “linearithmic” 😞
- $O(N^c)$ = polinomic 😡
- $O(c^N)$ = exponential 🤯
- $O(N!)$ = factorial 🤯

N : size of the problem, **c** : constant > 1

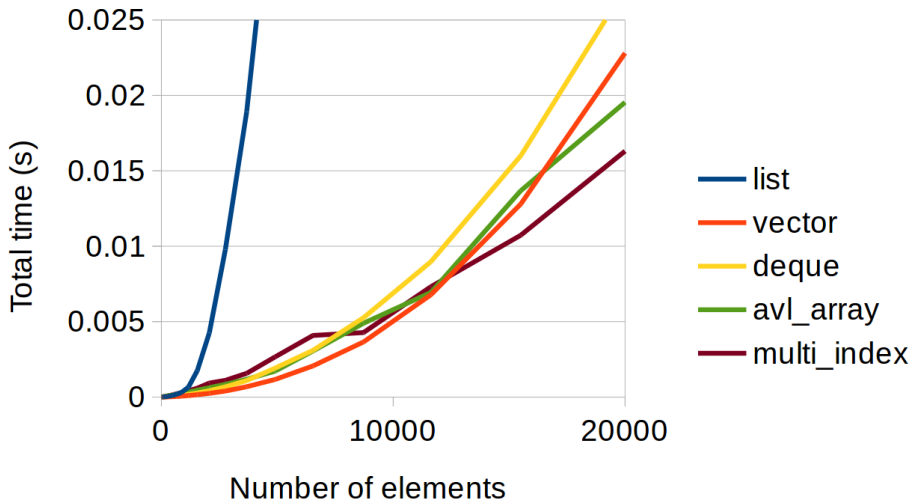
Computational complexity

	Random access	Insertion/ Extraction	Sum of both
Array			
List			
Super Tree			

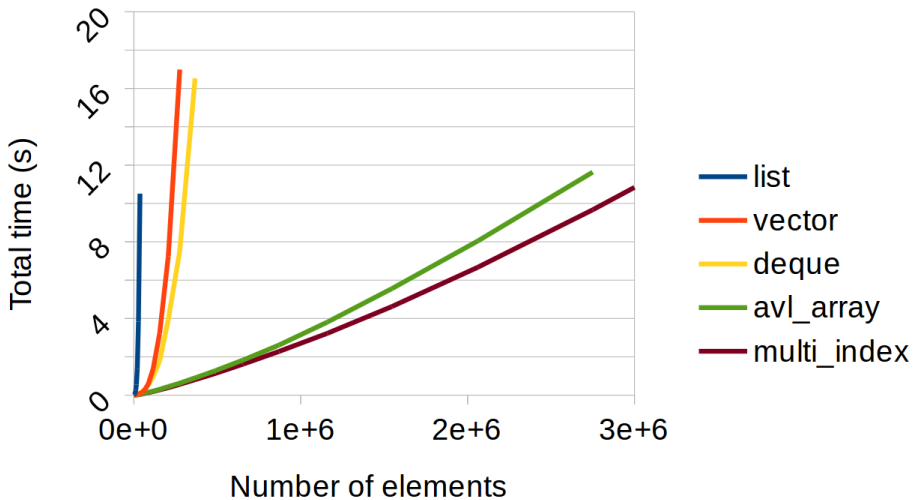
Computational complexity

	(1 rand. access + 1 ins./extr.) $\times N = \mathbf{total}$	
Array		
List		
Super Tree		

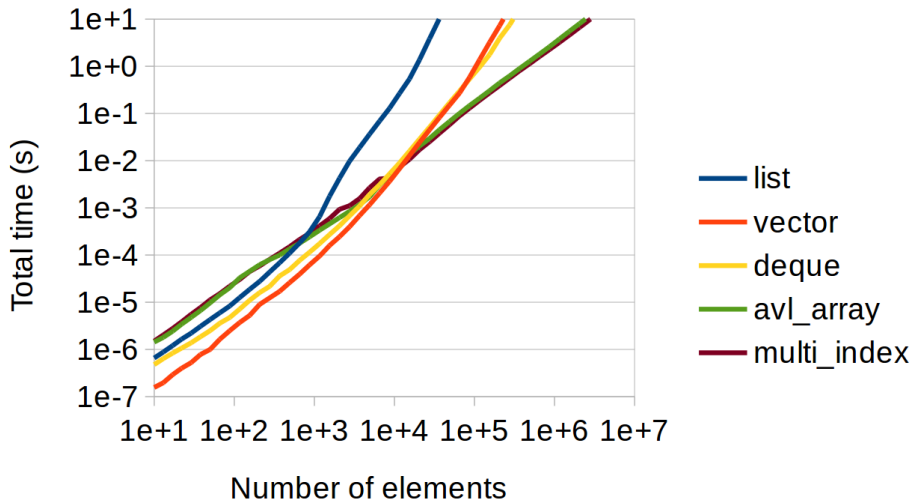
Results (1/3) — few elements



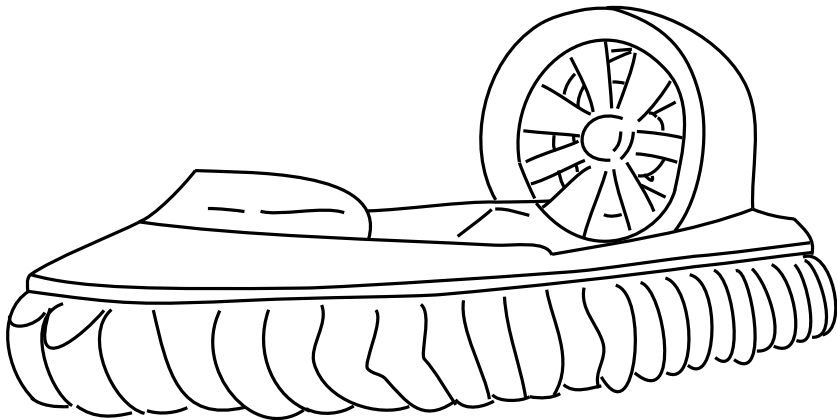
Results (2/3) — many elements



Results (3/3) — logarithmic scale



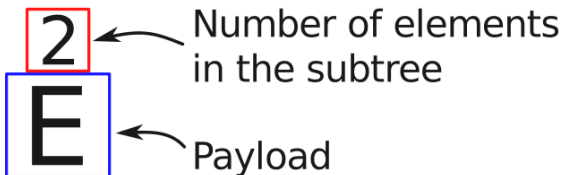
Ideal for the beach



Non proportional view

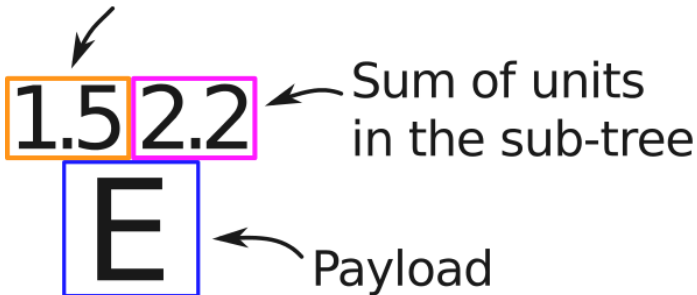
Legend in proportional view

1 element = 1 unit

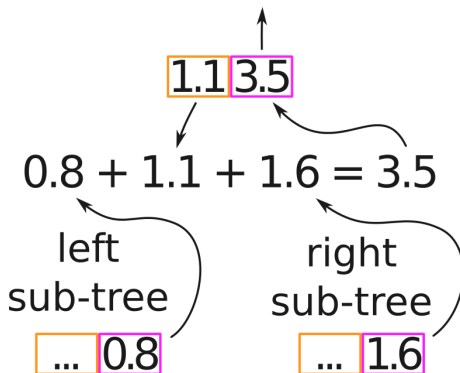


Legend in **non** proportional view

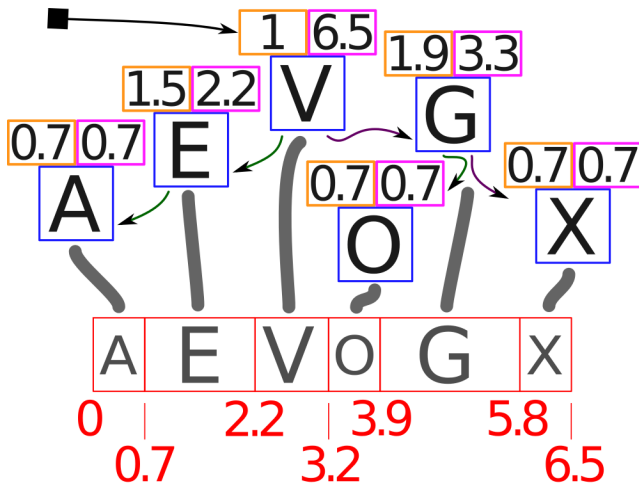
this element = 1.5 units



Sum in **non** proportional view



Non proportional view



Applications

Text editor

Sequence of lines

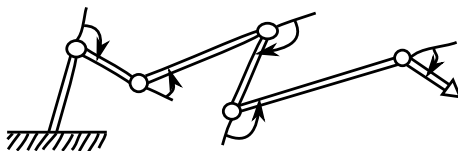
- Number of bytes
- Number of lines after word wrap
- Number of characters
- If not plain text, number of pixels

gtk

“Ad hoc” B+ tree with number of characters and lines

Robot arm or chain of molecules

- Sequence of traslation and rotation transformations
- **Non** proportional view operation:
matrix sum and product



Disk version: `shiftable_files`

- Implementation based on memory mapped files
- Horrible code (macros!)
- Metadata contained in the same file
- At closing time, choose:
 - 1 Recompact the file, or...
 - 2 leave it as is, with the metadata

How to keep track of the sections?

Using an in-memory sequence with **non** proportional view

Editing giant XML files

- A first pass can build an in-memory index (not necessarily complete)
- You can insert/extract nodes without rewriting the whole file
- You must keep the index updated, of course
- Recompact at closing?
 - 1 Yes: it becomes a normal XML again
 - 2 No: faster

Similar proposals

Multi Index (1/2)

```
boost::multi_index_container
<
  T,
  boost::multi_index::indexed_by
  <
    boost::multi_index::ranked_non_unique
    <
      boost::multi_index::identity<T>,
      unordered_less<T>
    >
  >
>
>
```

Multi Index (2/2)

```
template<typename T>
struct unordered_less
{
    bool operator() (const T &,
                    const T &) const
    {
        return false;
    }
};
```


Similar proposals in Boost (1/2)

- 2004 – The oldest mention (I don't know if implemented), by Peter Palotas

<http://lists.boost.org/Archives/boost/2004/03/62823.php>

- 2006 – “Hierarchical Data Structures” by Bernhard Reiter and René Rivera

<http://www.open-std.org/jtc1/sc22/wg21/docs/papers/2006/n2101.html#tr.hierarchy.augment>

- 2006 – “AVL Array” (horrible name, I know)

<http://sourceforge.net/projects/avl-array>

“Rank List” after debate in Boost forum

Similar proposals in Boost (2/2)

- 2012 – Countertree by Vadim Stadnik
[http://dl.dropbox.com/u/8437476/works/
countertree/doc/index.html](http://dl.dropbox.com/u/8437476/works/countertree/doc/index.html) (broken link)
- 2015 – SegmentedTree by Chris Clearwater
https://det.github.io/segmented_tree/

Similar proposals not in Boost

- “Simon Tatham’s Algorithms Page”

<https://www.chiark.greenend.org.uk/>

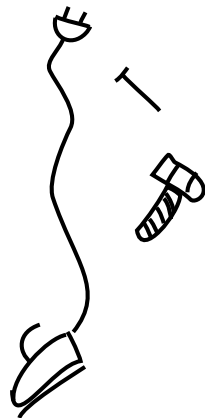
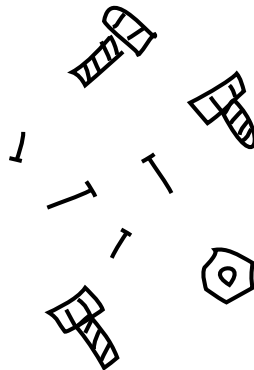
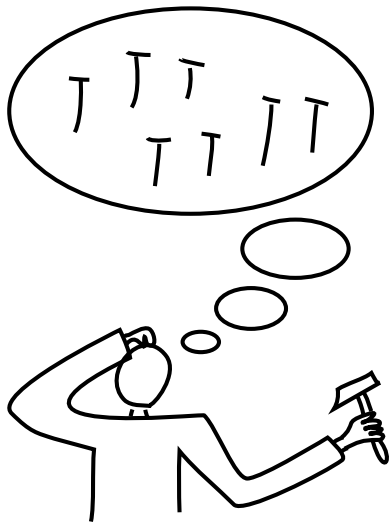
[~sgtatham/algorithms/cbtree.html](https://www.chiark.greenend.org.uk/~sgtatham/algorithms/cbtree.html)

“Counted B-trees: An enhancement to the well known B-tree algorithms to allow you to **look up items in the tree by numeric index**, or to **find the numeric index of an item**. Useful for finding percentiles, [...]”

Similar proposals in Python

- <https://pypi.python.org/pypi/rbtree>
- <https://pypi.python.org/pypi/pyavl>
- <https://pypi.python.org/pypi/blist>

Let's think about it



Thanks a lot

Any questions?