# Advanced Data Structures 











## Insertion Order Matters

- Suppose we create a BST of numbers in this order:

$$
4,2,1,3,6,5,7
$$

4


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$$
1,2,3,4,5,6,7
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## Tree Terminology

- The height of a tree is the number of nodes in the longest path from the root to a leaf.



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## Keeping the Height Low

- Almost all BST operations have time complexity based on height:
- Insertion: O(h)
- Search: O(h)
- Deletion: O(h)
- Keeping the height low will make these operations much more efficient.
- How do we do this?


## Tree Rotations

- One common way of keeping tree heights low is to reshape the BST when it gets too high.
- One way to accomplish this is a tree rotation, which locally rearranges nodes.


## Tree Rotations




## $2$



, 2
‘


5










## Let's Code it Up!

## When to Rotate?

- The actual code for rotations is not too complex.
- Deciding when and where to rotate the tree, on the other hand, is a bit involved.
- There are many schemes we can use to determine this:
- AVL trees maintain balance information in each node, then rotate when the balance is off.
- Red/Black trees assign each node a color, then rotate when certain color combinations occur.


## An Interesting Observation

## Random Binary Search Trees

- If we build a binary search tree with totally random values, the resulting tree is (with high probability) within a constant factor of balanced.
- Approximately $4.3 \ln n$
- Moreover, the average depth of a given node is often very low.
- Approximately $2 \ln n$.
- If we structure the BST as if it were a random tree, we get (with high probability) a very good data structure!


## Treaps

- A treap is a data structure that combines a binary search tree and a binary heap.
- Each node stores two pieces of information:
- The piece of information that we actually want to store, and
- A random real number.
- The tree is stored such that
- The nodes are a binary search tree when looking up the information, and
- The nodes are a binary heap with respect to the random real number.



## Treaps are Wonderful

- With very high probability, the height of an $n$-node treap is $\mathrm{O}(\log n)$.
- Insertion is surprisingly simple once we have code for tree rotations.
- Deletion is straightforward once we have code for tree rotations.


## Inserting into a Treap

- Insertion into a treap is a combination of normal BST insertion and heap insertion.
- First, insert the node doing a normal BST insertion. This places the value into the right place.
- Next, bubble the node upward in the tree by rotating it with its parent until its value is smaller than its parent.







## Let's Code it Up!

## Removing from a Treap

- In general, removing a node from a BST is quite difficult because we have to make sure not to lose any nodes.
- For example, how do you remove the root of this tree?

- However, removing leaves is very easy, since they have no children.


## Removing from a Treap

- It would seem that, since a treap has extra structure on top of that of a BST, that removing from a treap would be extremely hard.
- However, it's actually quite simple:
- Keep rotating the node to delete with its larger child until it becomes a leaf.
- Once the node is a leaf, delete it.







## Summary of Treaps

- Treaps give a (reasonably) straightforward way to guarantee that the height of a BST is not too great.
- Insertion into a treap is similar to insertion into a BST followed by insertion into a binary heap.
- Deletion from a treap is similar to the bubble-down step from a heap.
- All operations run in expected $\mathrm{O}(\log n)$ time.


## A Survey of Other Data Structures

## Data Structures so Far

- We have seen many data structures over the past few weeks:
- Dynamic arrays.
- Linked lists.
- Hash tables.
- Tries.
- Binary search trees (and treaps).
- Binary heaps.
- These are the most-commonly-used data structures for general data storage.


## Specialized Data Structures

- For applications that manipulate specific types of data, other data structures exist that make certain operations surprisingly fast and efficient.
- Many critical applications of computers would be impossible without these data structures.


## k-d Trees

Suppose that you want to efficiently store points in $k$-dimensional space.

How might you organize the data to efficiently query for points within a region?


## Search for (1, 4, 4)

$(3,1,4)$


$$
\begin{gathered}
\mathbf{V} \\
(4,0,6)
\end{gathered}
$$



「
$(7,1,6)$

## Search for (1, 4, 4)

$(3,1,4)$


$$
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\mathbf{V} \\
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「
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$$
\begin{gathered}
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(4, \mathbf{0}, 6)
\end{gathered}
$$



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「
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## The Intuition

## The Intuition



## The Intuition



## The Intuition



## The Intuition



## The Intuition



Values less than two
Values greater than two

## The Intuition



Values less than two
Values greater than two

## Key Idea: Split Space in Half





Upper half-space


Lower half-space










Nearest-Neighbor Lookup






$y=y_{0}$

$y=y_{0}$

$y=y_{0}$


## k-d Trees

- Assuming the points are nicely distributed, nearest-neighbor searches in $k$-d trees can run faster than $\mathrm{O}(n)$ time.
- Applications in computational geometry (collision detection), machine learning (nearest-neighbor classification), and many other places.


## Suffix Trees

## String Processing

- In computational biology, strings are enormously useful for storing DNA and RNA.
- Many important questions in biology can be addressed through string processing:
- What is the most plausible evolutionary history of the following genomes?
- Are there particular gene sequences that appear with high frequency within a genome?


## Suffix Trees

- A suffix tree is a (slightly modified) trie that stores all suffixes of a string $S$.
- Here is the suffix tree for "dikdik;" the \$ is a marker for "end-of-string."



## Suffix Trees

- Important, nontrivial, nonobvious fact: A suffix tree for a string of $n$ characters can be built in time $O(n)$.
- Given a string of length $m$, we can determine whether it is a substring of the original string in time $\mathrm{O}(m)$.



## Suffix Trees

- Other applications of suffix trees:
- Searching for one genome within another allowing for errors, insertions, and deletions in time $\mathrm{O}(n+m)$.
- Finding the longest common substring of two sequences in time $\mathrm{O}(n+m)$.
- Improving the performance of data compression routines by finding long repeated strings efficiently.


## Bloom Filters

## Distributing Data

- Websites like Google and Facebook deal with enormous amounts of data.
- Probably measured in hundreds of millions of gigabytes (hundreds of petabytes).
- There is absolutely no way to store this on one computer.
- Instead, data must be stored on multiple computers networked together.


## Looking up Data

- Suppose you are at Google implementing search.
- When you get a search query, you have to be able to know which computer knows what pages to display for that query.
- Network latency is, say, 2 ms between you and each computer.
- If you have one thousand computers to search, you can't just query each one and ask.


## Bloom Filters

- A Bloom filter is a data structure similar to a set backed by a hash table.
- Stores a set of values in a way that may lead to false positives:
- If the Bloom filter says that an object is not present, it is definitely not present.
- If the Bloom filter says that an object is present, it may actually not be present.


## Bloom Filters

## $\begin{array}{llllllllllllll}\mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N}\end{array}$

## Bloom Filters

Value<br>One

## $\begin{array}{llllllllllllll}\mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N} & \mathbf{N}\end{array}$

## Bloom Filters

## Value <br> One



## Bloom Filters

## Value <br> One



## Bloom Filters

## $\mathbf{N} \mathbf{N} \quad \mathbf{N} \quad \mathbf{Y} \mathbf{N} \mathbf{N} \mathbf{Y} \mathbf{N} \mathbf{Y} \mathbf{N} \mathbf{N} \mathbf{Y} \mathbf{N} \mathbf{Y}$

## Bloom Filters

Value<br>Two

## $\mathbf{N} \mathbf{N} \quad \mathbf{N} \quad \mathbf{Y} \mathbf{N} \mathbf{N} \mathbf{Y} \mathbf{N} \mathbf{Y} \mathbf{N} \mathbf{N} \mathbf{Y} \mathbf{N} \mathbf{Y}$

## Bloom Filters

Value<br>Two



## Bloom Filters

Value<br>Two



## Bloom Filters

## $\begin{array}{llllllllllllll}\mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y}\end{array}$

## Bloom Filters

Value<br>One

| $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |

## Bloom Filters

Value<br>One



## Bloom Filters

## $\begin{array}{llllllllllllll}\mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y}\end{array}$

## Bloom Filters

Value<br>Two

| $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |

## Bloom Filters

Value<br>Two



## Bloom Filters

## $\begin{array}{llllllllllllll}\mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y}\end{array}$

## Bloom Filters

Value Three

| $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{N}$ | $\mathbf{Y}$ | $\mathbf{N}$ | $\mathbf{Y}$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |

## Bloom Filters

Value<br>Three



## Bloom Filters

## $\begin{array}{llllllllllllll}\mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y}\end{array}$

## Bloom Filters

## Value

Four

## $\begin{array}{llllllllllllll}\mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{Y} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{N} & \mathbf{Y} & \mathbf{N} & \mathbf{Y}\end{array}$

## Bloom Filters

Value<br>Four



## Bloom Filters and Networks

- Bloom filters can be used to mitigate the networking problem from earlier.
- Have each computer store a Bloom filter of what's stored on each other computer.
- To determine which computer has some data:
- Look up that value in each Bloom filter.
- Call up just the computers that might have it.
- Since Bloom filter lookup is substantially faster than a network query (probably 1000-10,000x), this solution is used extensively in practice.

Data structures make it possible to solve important problems at scale.

You get to decide which problems we'll be using them for.

