



2/26/08

## • Data-structure augmentation

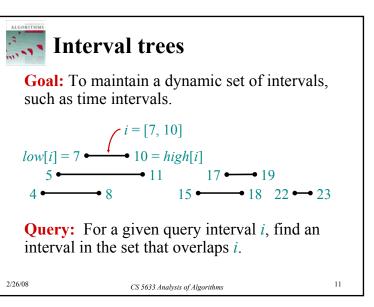
## Methodology: (e.g., order-statistics trees)

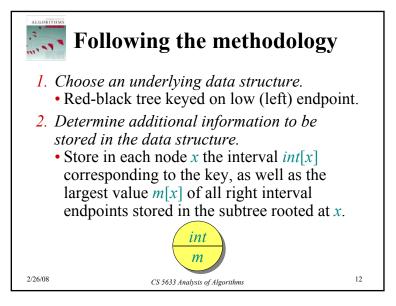
- 1. Choose an underlying data structure (*red-black trees*).
- 2. Determine additional information to be stored in the data structure (*subtree sizes*).
- 3. Verify that this information can be maintained for modifying operations (RB-INSERT, RB-DELETE *don't forget rotations*).
- 4. Develop new dynamic-set operations that use the information (OS-SELECT *and* OS-RANK).

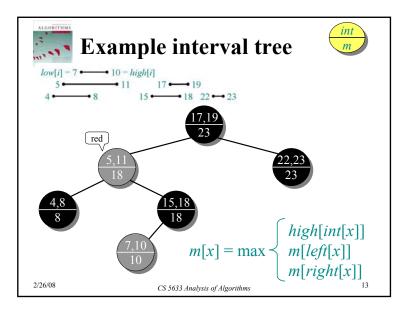
These steps are guidelines, not rigid rules.

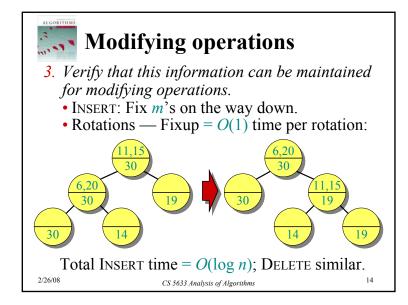
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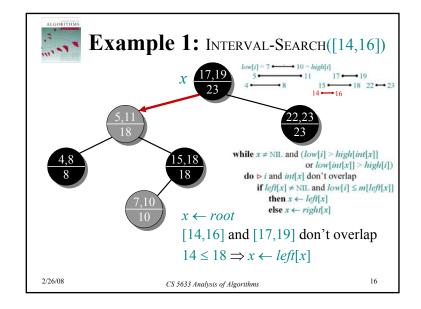


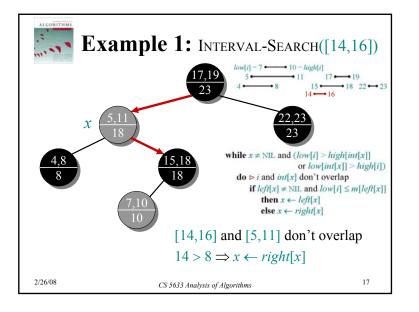


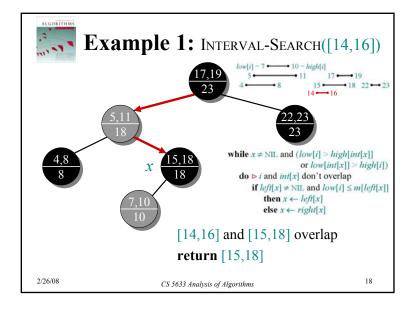


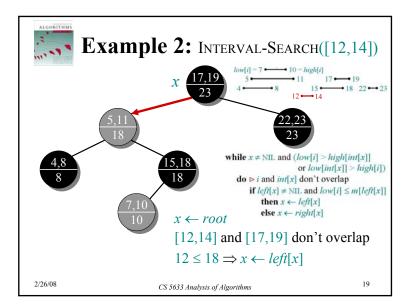


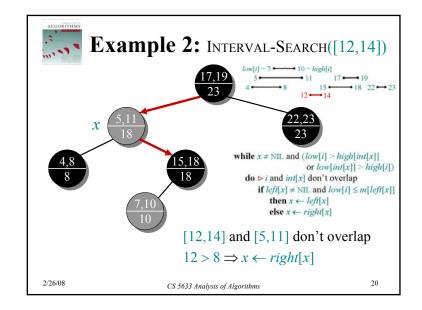


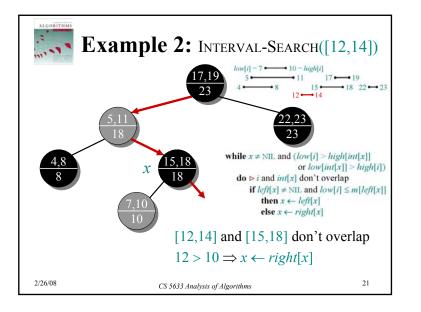


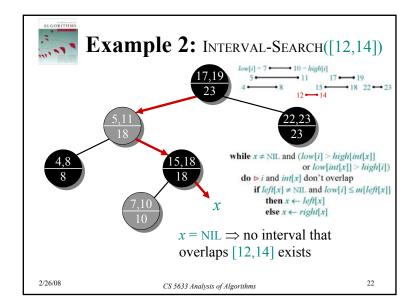


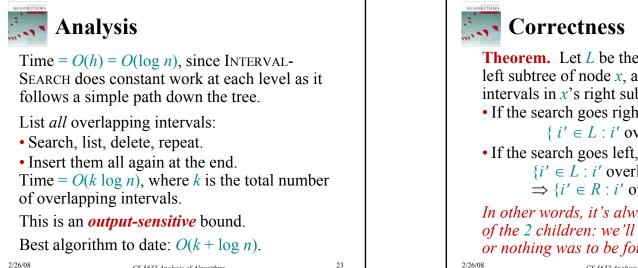












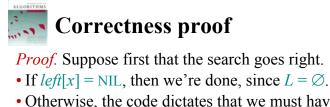
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**Theorem.** Let *L* be the set of intervals in the left subtree of node x, and let R be the set of intervals in x's right subtree.

• If the search goes right, then

 $\{i' \in L : i' \text{ overlaps } i\} = \emptyset.$ • If the search goes left, then  $\{i' \in L : i' \text{ overlaps } i\} = \emptyset$  $\Rightarrow$  {*i*'  $\in$  *R* : *i*' overlaps *i* } =  $\emptyset$ .

In other words, it's always safe to take only 1 of the 2 children: we'll either find something, or nothing was to be found.



• Otherwise, the code dictates that we must have low[i] > m[left[x]]. The value m[left[x]] corresponds to the right endpoint of some interval  $j \in L$ , and no other interval in *L* can have a larger right endpoint than high(j).

$$i$$

$$high(j) = m[left[x]] \qquad i$$

$$low(i)$$
• Therefore,  $\{i' \in L : i' \text{ overlaps } i\} = \emptyset$ .
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