Part III

Data Structures



Abstract Data Type

An abstract data type (ADT) is defined by an interface of operations or methods that can be performed and that have a defined behavior.

The data types in this lecture all operate on objects that are represented by a [key, value] pair.

- The key comes from a totally ordered set, and we assume that there is an efficient comparison function.
- The value can be anything; it usually carries satellite information important for the application that uses the ADT.

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- S. successor(x): Return pointer to the next larger element in S or null if x is maximum.

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- ► *S*. concatenate(S'): $S := S \cup S'$. Requires key[S. maximum()] \leq key[S'. minimum()].
- ► *S.* decrease-key(x, k): Replace key[x] by $k \le key[x]$.

Examples of ADTs

Stack:

- \triangleright S. push(x): Insert an element.
- ▶ *S.* pop(): Return the element from *S* that was inserted most recently; delete it from *S*.
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Priority-Queue:

- S. insert(x): Insert an element.
- S. delete-min(): Return the element with lowest key-value; delete it from S.

7 Dictionary

Dictionary:

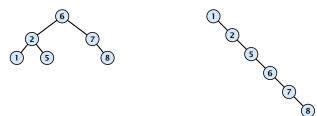
- S. insert(x): Insert an element x.
- **S.** delete(x): Delete the element pointed to by x.
- ▶ *S.* search(k): Return a pointer to an element e with key[e] = k in S if it exists; otherwise return null.

7.1 Binary Search Trees

An (internal) binary search tree stores the elements in a binary tree. Each tree-node corresponds to an element. All elements in the left sub-tree of a node v have a smaller key-value than $\ker[v]$ and elements in the right sub-tree have a larger-key value. We assume that all key-values are different.

(External Search Trees store objects only at leaf-vertices)

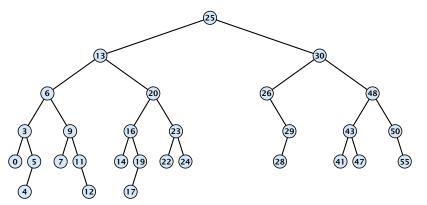
Examples:



7.1 Binary Search Trees

We consider the following operations on binary search trees. Note that this is a super-set of the dictionary-operations.

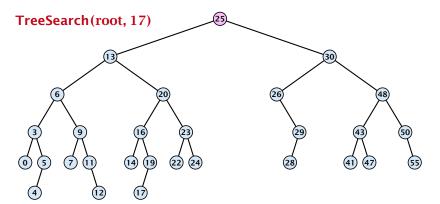
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- 1: **if** x = null or k = key[x] **return** x
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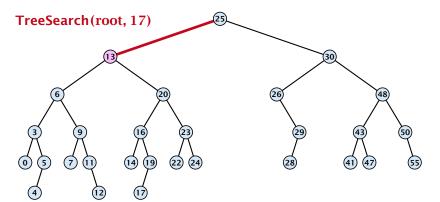


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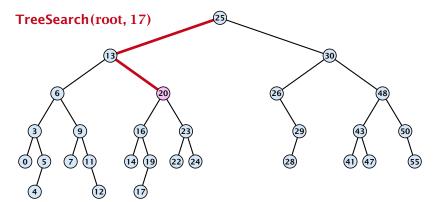




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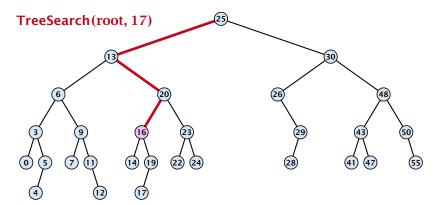


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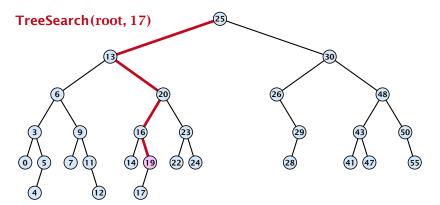


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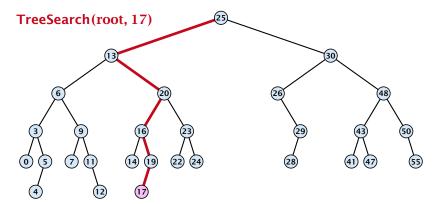


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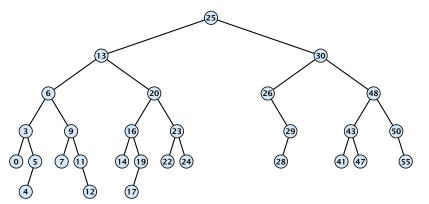


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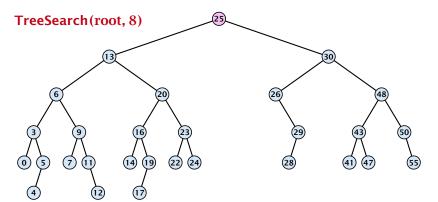


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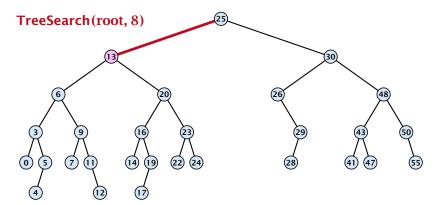




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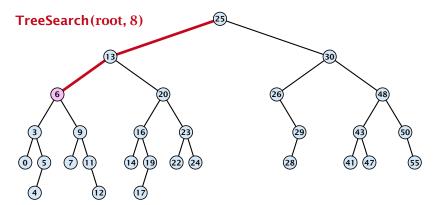




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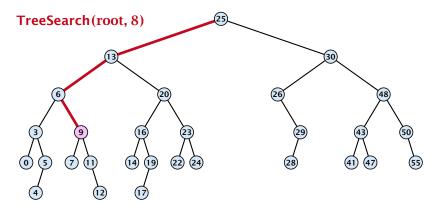


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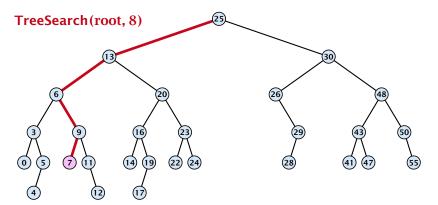




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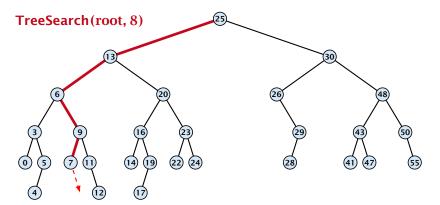


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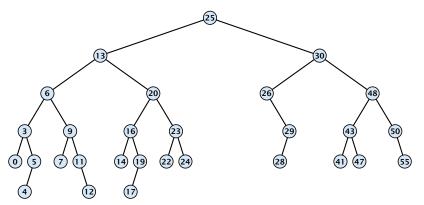
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Binary Search Trees: Minimum

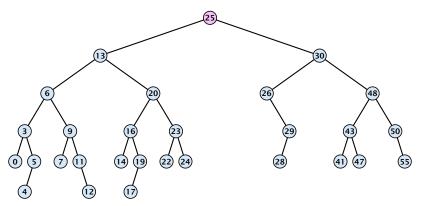


Algorithm 2 TreeMin(x)

- 1: **if** x = null or left[x] = null return x
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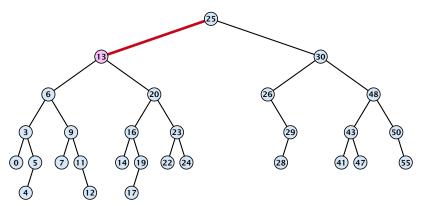
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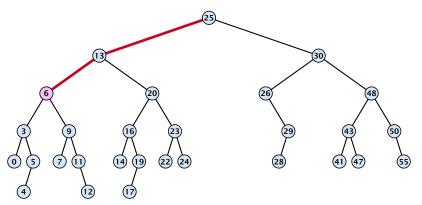
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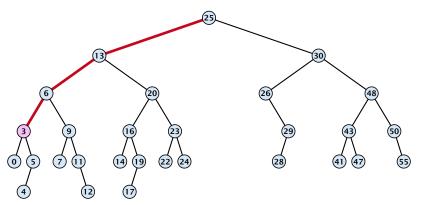
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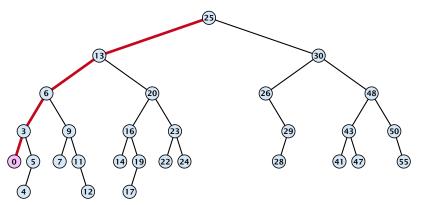
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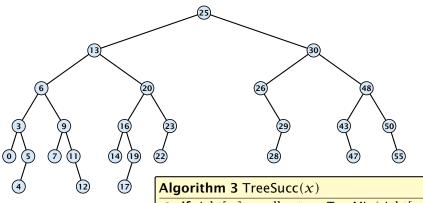
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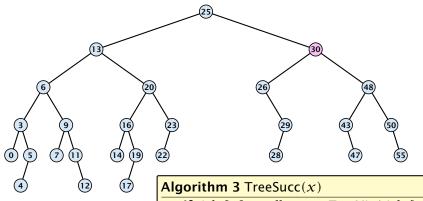
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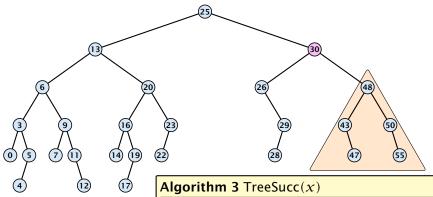
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- 5: **return** y;





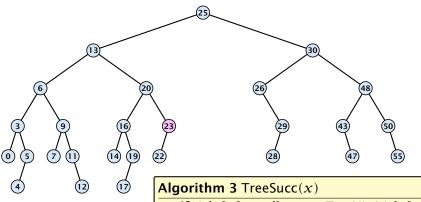
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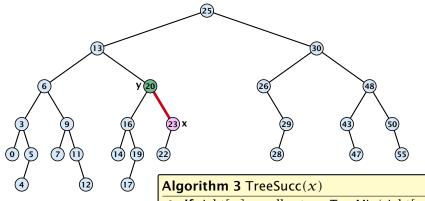
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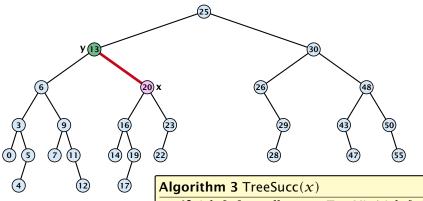
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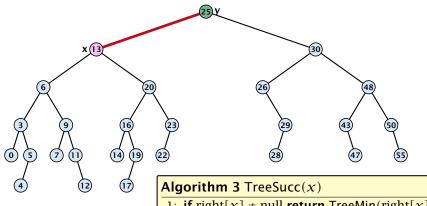
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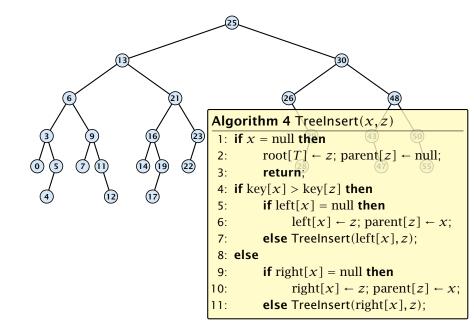
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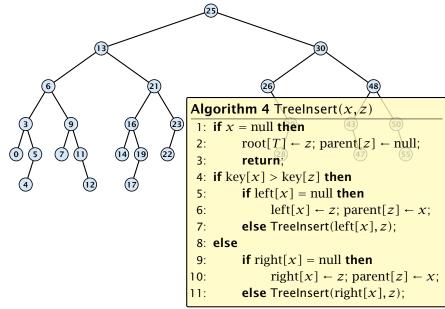


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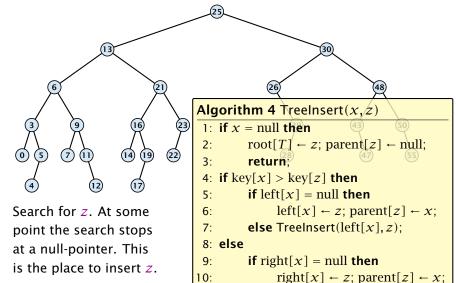




Insert element not in the tree.



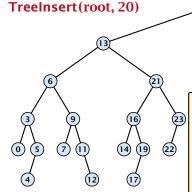
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11:

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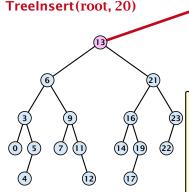
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Search for z. At some point the search stops at a null-pointer. This is the place to insert z.

- 1: if x = null then2: $\text{root}[T] \leftarrow z$; parent $[z] \leftarrow \text{null}$;
 - 3: return
- 4: **if** key[x] > key[z] **then**
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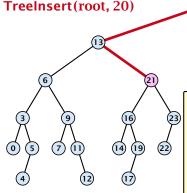
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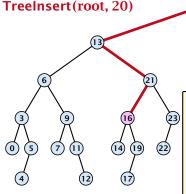
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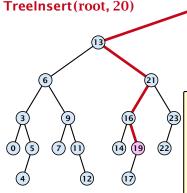
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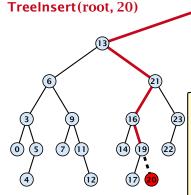
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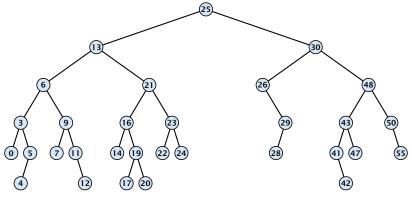
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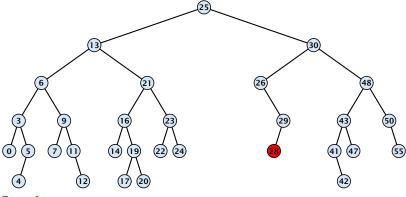
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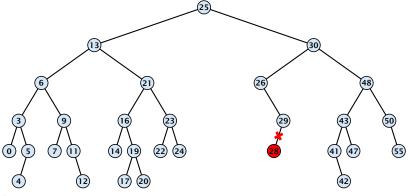




Case 1:

Element does not have any children

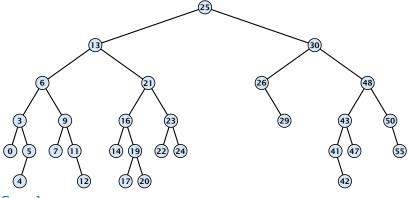
Simply go to the parent and set the corresponding pointer to null.



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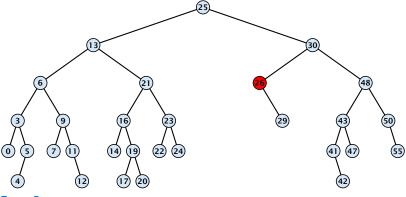
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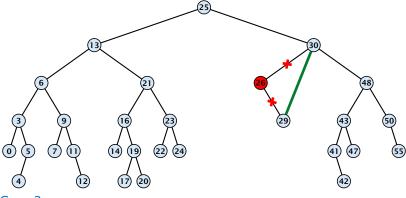
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Case 2:

Element has exactly one child

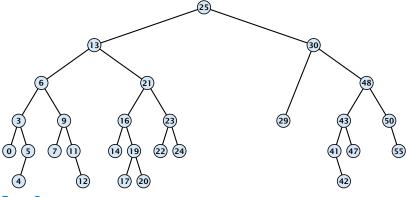
Splice the element out of the tree by connecting its parent to its successor.



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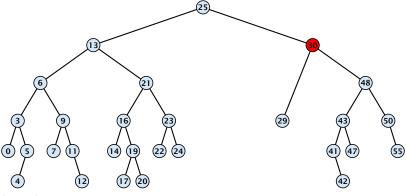
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Case 2:

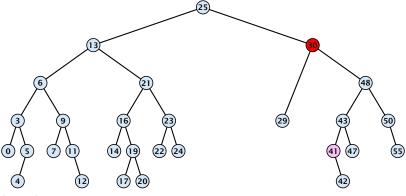
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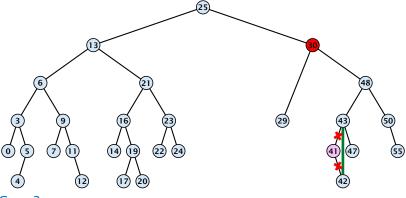
Case 3:

- Find the successor of the element
- Splice successor out of the tree
- Replace content of element by content of successor



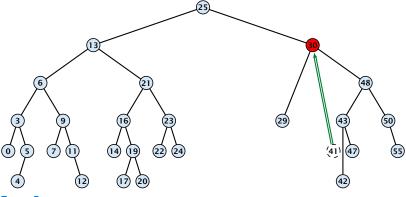
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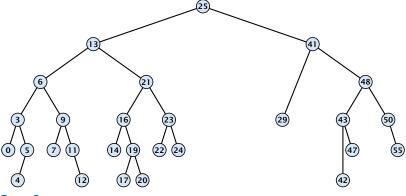
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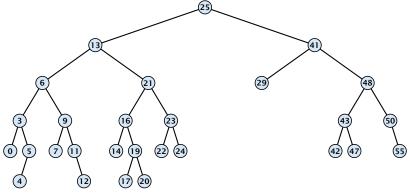
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```
Algorithm 9 TreeDelete(z)
 1: if left[z] = null or right[z] = null
          then \gamma \leftarrow z else \gamma \leftarrow \text{TreeSucc}(z); select \gamma to splice out
 3: if left[\gamma] \neq null
         then x \leftarrow \text{left}[y] else x \leftarrow \text{right}[y]; x is child of y (or null)
 5: if x \neq \text{null then parent}[x] \leftarrow \text{parent}[y]; parent[x] is correct
 6: if parent[\gamma] = null then
 7: root[T] \leftarrow x
 8: else
 9: if \gamma = \text{left[parent}[\gamma]] then
                                                                  fix pointer to x
10:
                left[parent[v]] \leftarrow x
    else
11:
12.
        right[parent[y]] \leftarrow x
13: if y \neq z then copy y-data to z
```

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Balanced Binary Search Trees

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With each insert- and delete-operation perform local adjustments to guarantee a height of $\mathcal{O}(\log n)$.

AVL-trees, Red-black trees, Scapegoat trees, 2-3 trees, B-trees, AA trees, Treaps

similar: SPLAY trees.



Definition 1

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A red black tree is a balanced binary search tree in which each internal node has two children. Each internal node has a color, such that

1. The root is black.

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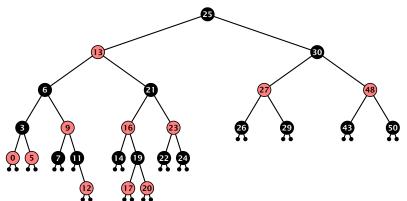
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Red Black Trees: Example



Lemma 2

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The black height $\mathrm{bh}(v)$ of a node v in a red black tree is the number of black nodes on a path from v to a leaf vertex (not counting v).

Lemma 2

A red-black tree with n internal nodes has height at most $\mathcal{O}(\log n)$.

Definition 3

The black height bh(v) of a node v in a red black tree is the number of black nodes on a path from v to a leaf vertex (not counting v).

We first show:

Lemma 4

A sub-tree of black height bh(v) in a red black tree contains at least $2^{bh(v)}-1$ internal vertices.

Proof of Lemma 4.

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Induction on the height of v.

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- ▶ The black height of v is 0.
- ► The sub-tree rooted at v contains $0 = 2^{bh(v)} 1$ inner vertices.

Proof (cont.)

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induction step

Supose v is a node with height(v) > 0.

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- **By** induction hypothesis both sub-trees contain at least $2^{\text{bh}(v)-1}-1$ internal vertices.
- ► Then T_v contains at least $2(2^{\text{bh}(v)-1}-1)+1 \ge 2^{\text{bh}(v)}-1$ vertices.



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Hence, $h \le 2\log(n+1) = \mathcal{O}(\log n)$.



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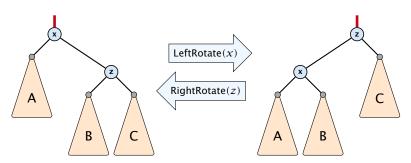
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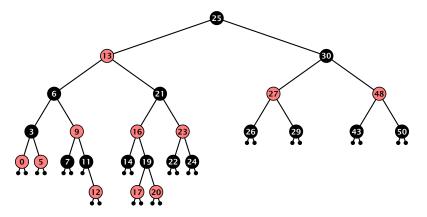
We need to adapt the insert and delete operations so that the red black properties are maintained.

Rotations

The properties will be maintained through rotations:



Red Black Trees: Insert

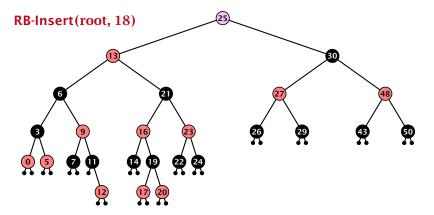


Insert:

- first make a normal insert into a binary search tree
- then fix red-black properties



Red Black Trees: Insert

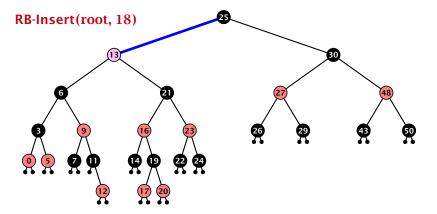


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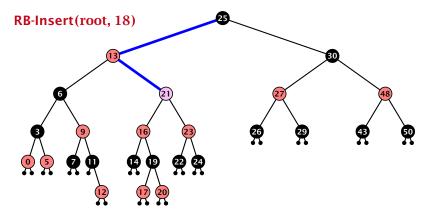
Red Black Trees: Insert



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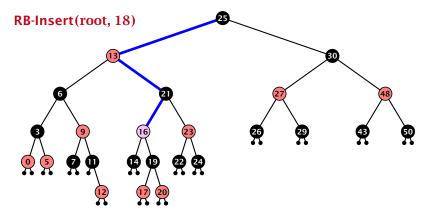
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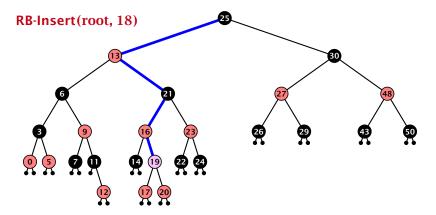
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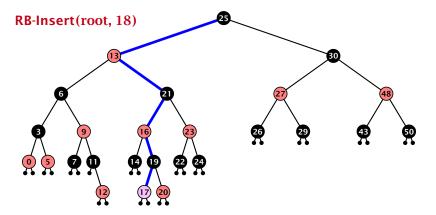
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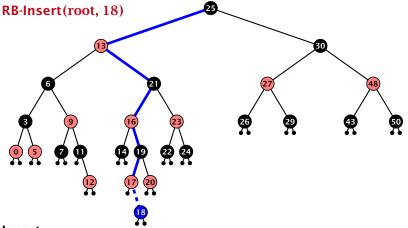
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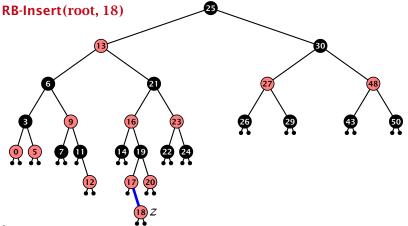
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 - either both of them are red (most important case)
 - or the parent does not exist (violation since root must be black)

If z has a parent but no grand-parent we could simply color the parent/root black; however this case never happens.

```
Algorithm 10 InsertFix(z)
 1: while parent[z] \neq null and col[parent[z]] = red do
         if parent[z] = left[gp[z]] then
 2:
 3:
               uncle \leftarrow right[grandparent[z]]
               if col[uncle] = red then
 4:
                    col[p[z]] \leftarrow black; col[u] \leftarrow black;
 5:
                    col[gp[z]] \leftarrow red; z \leftarrow grandparent[z];
 6:
 7:
              else
                    if z = right[parent[z]] then
 8:
                         z \leftarrow p[z]; LeftRotate(z);
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```

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 4:
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6. Feb. 2022

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                                                               Case 2: uncle black
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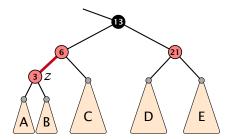


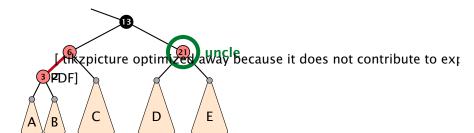
7.2 Red Black Trees 6. Feb. 2022

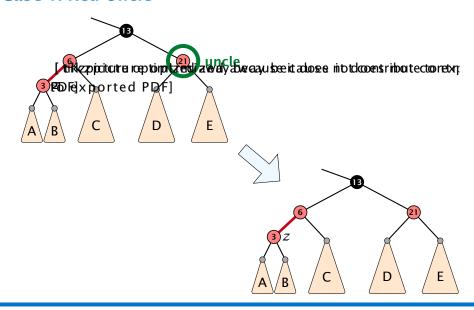
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                                                                   2a: z right child
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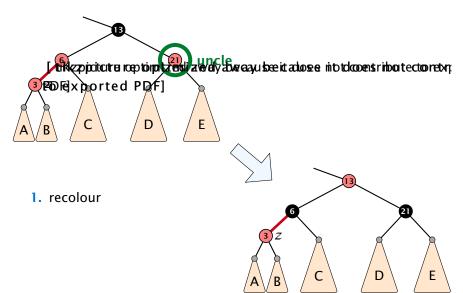
6. Feb. 2022 42/308

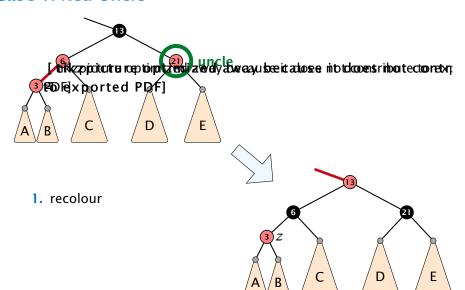
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```





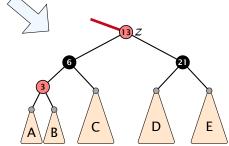








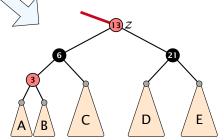
- 1. recolour
- 2. move z to grand-parent



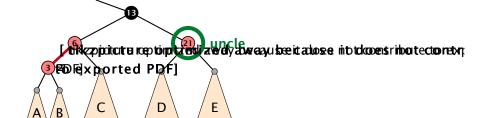
7.2 Red Black Trees



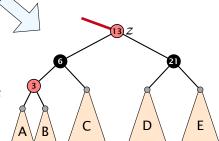
- 1. recolour
- 2. move z to grand-parent
- 3. invariant is fulfilled for new z

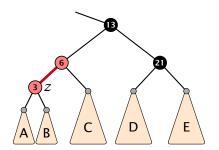


7.2 Red Black Trees

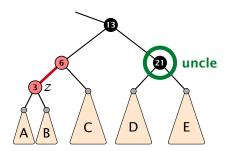


- recolour
- 2. move z to grand-parent
- 3. invariant is fulfilled for new z
- 4. you made progress





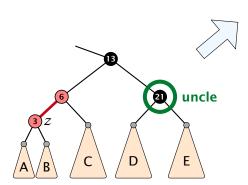
[tikzpicture optimized away because it does not contribute to exp PDF]



1. rotate around grandparent

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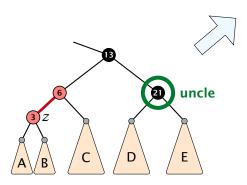
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1. rotate around grandparent

2. [reil@appointutenepropriatelolistwalyabaraayubeeit.adopee intotalooese intotaloo

Pholipskyhoerityehod pP10 pFerty holds





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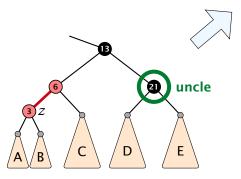
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1. rotate around grandparent

2. [reikzphoùchto repromptei eloliez verdy abverayu bee it actions e into toloroes inhout ector retyr.

Poliejsk phoeingend parto prety holds

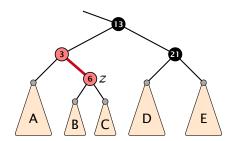
3. you have a red black tree



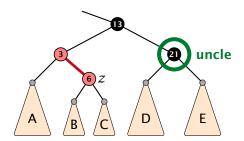


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Performing Case 1 at most $\mathcal{O}(\log n)$ times and every other case at most once, we get a red-black tree. Hence $\mathcal{O}(\log n)$ re-colorings and at most 2 rotations.

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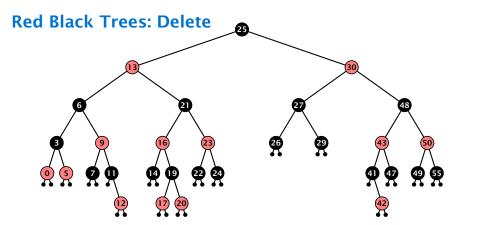
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- If you delete the root, the root may now be red.

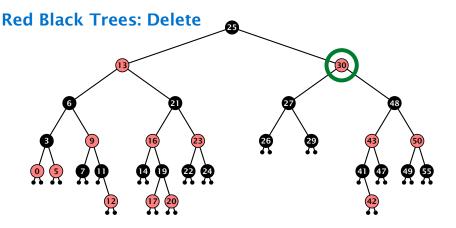
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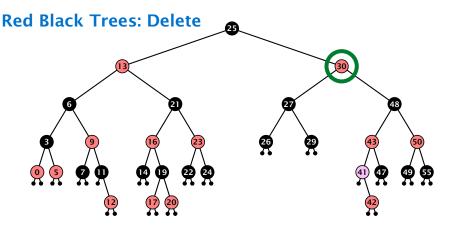
- Parent and child of x were red; two adjacent red vertices.
- If you delete the root, the root may now be red.
- Every path from an ancestor of x to a descendant leaf of x changes the number of black nodes. Black height property might be violated.





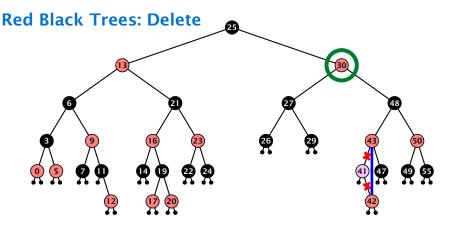
Case 3:

- do normal delete
- when replacing content by content of successor, don't change color of node



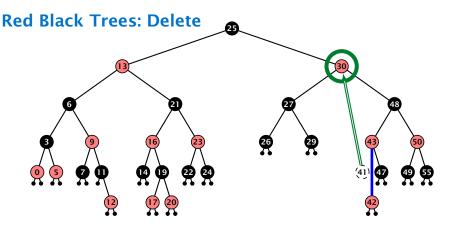
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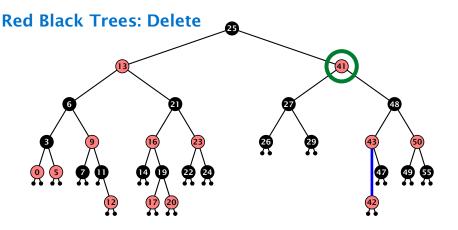
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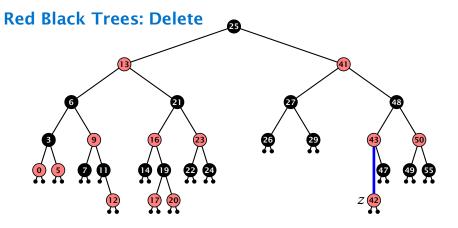
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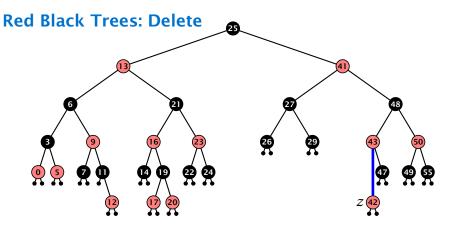
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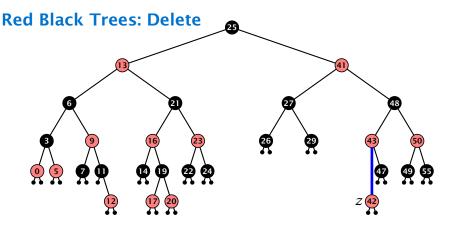
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deleting black node messes up black-height property



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Delete:

- deleting black node messes up black-height property
- ightharpoonup if z is red, we can simply color it black and everything is fine
- the problem is if z is black (e.g. a dummy-leaf); we call a fix-up procedure to fix the problem.

Invariant of the fix-up algorithm

► the node z is black

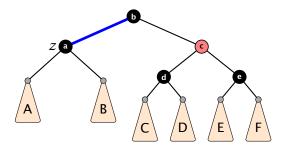
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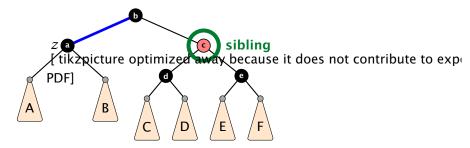
- ▶ the node *z* is black
- if we "assign" a fake black unit to the edge from z to its parent then the black-height property is fulfilled

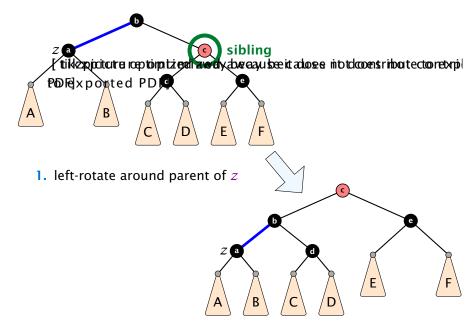
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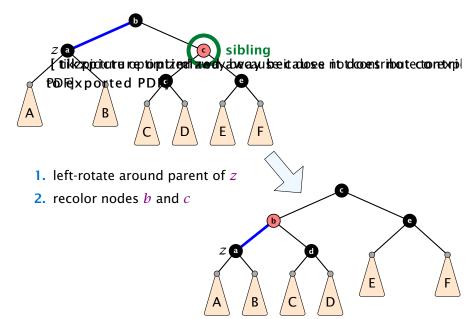
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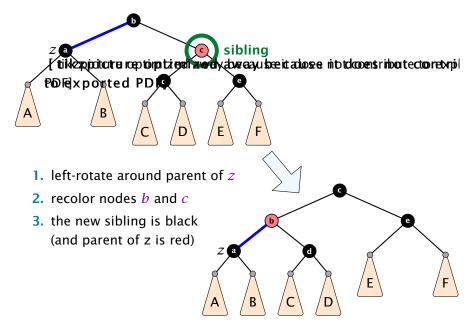
Goal: make rotations in such a way that you at some point can remove the fake black unit from the edge.

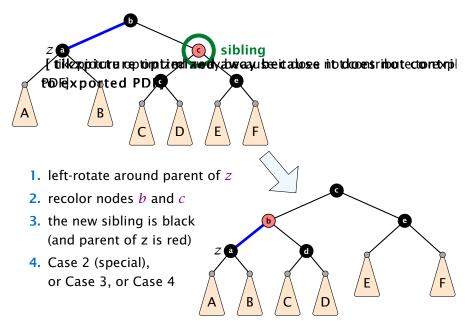


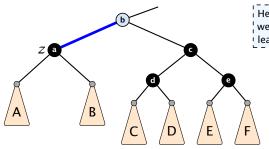




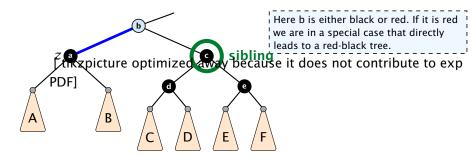


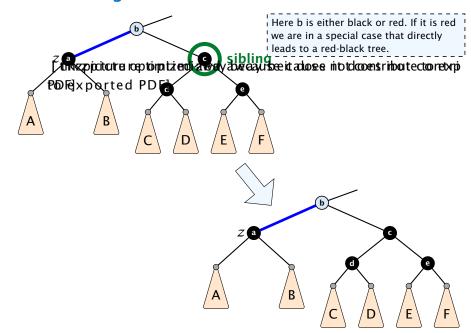


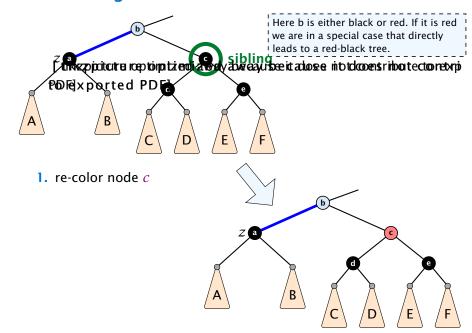


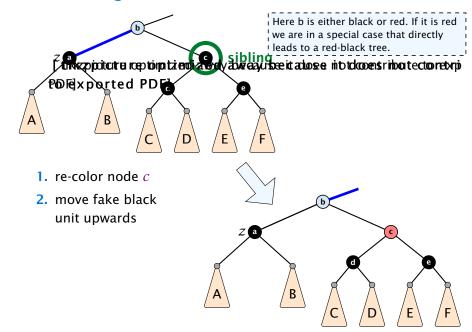


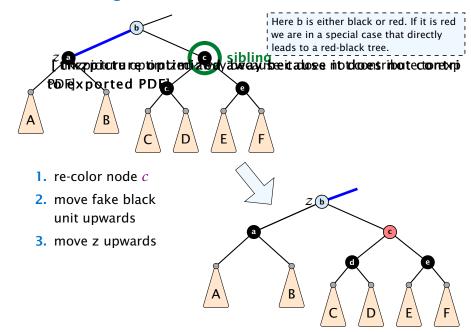
Here b is either black or red. If it is red we are in a special case that directly leads to a red-black tree.

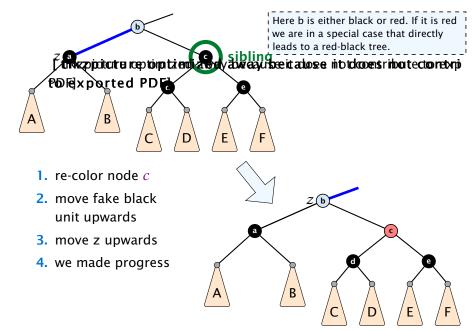




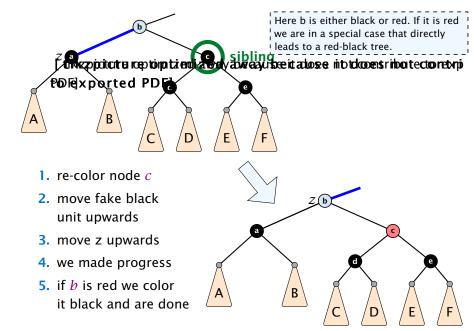


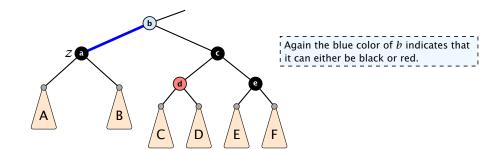




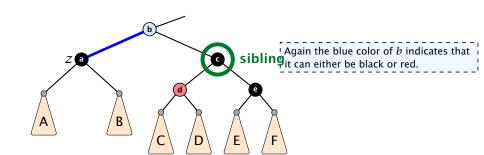


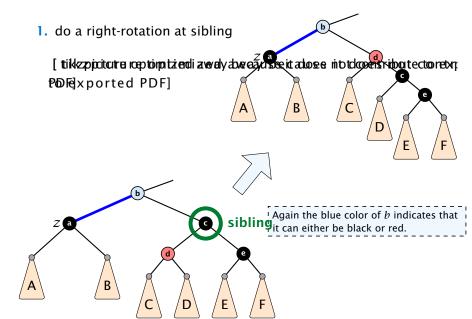
Case 2: Sibling is black with two black children

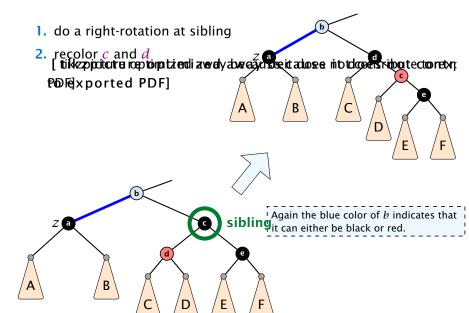


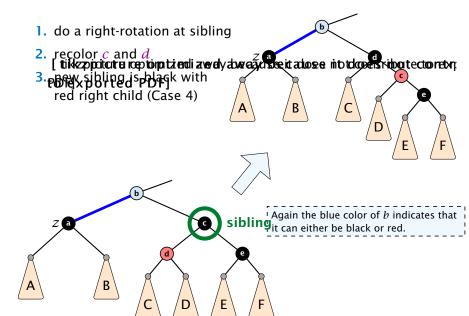


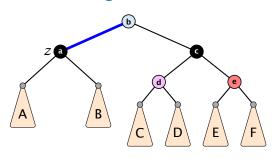
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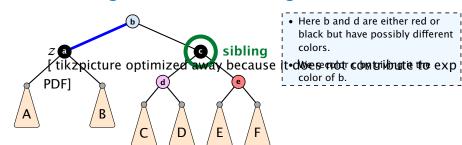


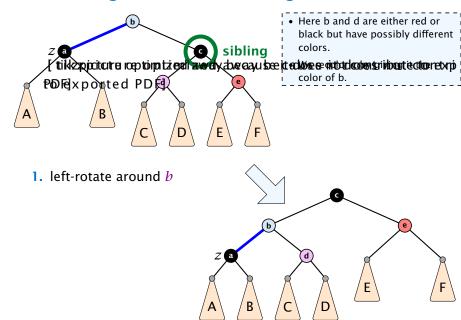


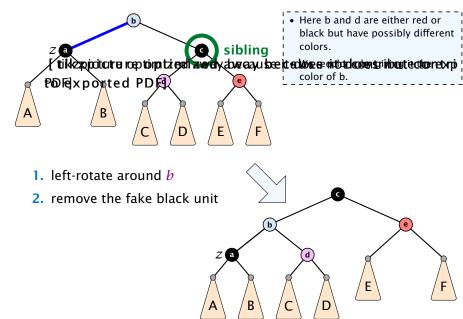


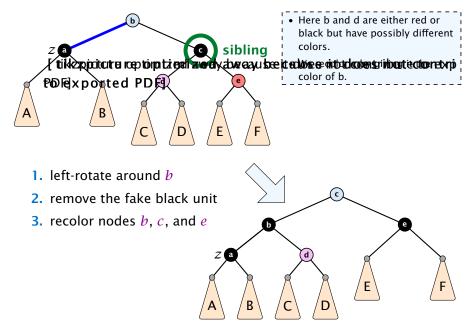


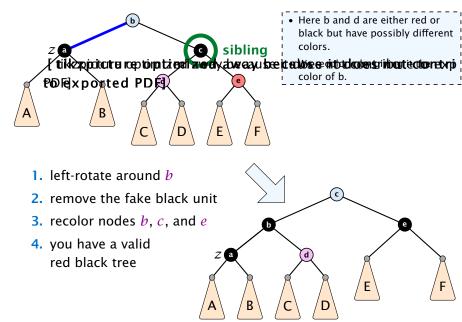
- Here b and d are either red or black but have possibly different colors.
- We recolor c by giving it the color of b.











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Performing Case 2 at most $\mathcal{O}(\log n)$ times and every other step at most once, we get a red black tree. Hence, $\mathcal{O}(\log n)$ re-colorings and at most 3 rotations.

Disadvantage of balanced search trees:

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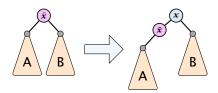
- + after access, an element is moved to the root; splay(x) repeated accesses are faster
- only amortized guarantee
- read-operations change the tree

find(x)

- search for x according to a search tree
- let \bar{x} be last element on search-path
- ightharpoonup splay (\bar{x})

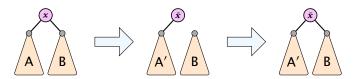
insert(x)

- search for x; \bar{x} is last visited element during search (successer or predecessor of x)
- splay(\bar{x}) moves \bar{x} to the root
- insert x as new root

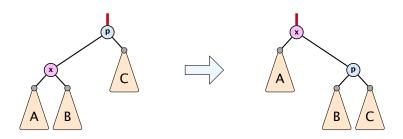


delete(x)

- search for x; splay(x); remove x
- **>** search largest element \bar{x} in A
- splay(\bar{x}) (on subtree A)
- connect root of B as right child of \bar{x}



Move to Root



How to bring element to root?

- one (bad) option: moveToRoot(x)
- iteratively do rotation around parent of x until x is root
- if x is left child do right rotation otw. left rotation

7.3 Splay Trees

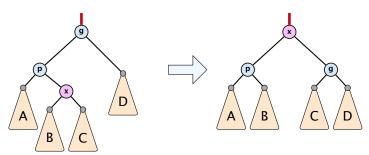
Splay: Zig Case



better option splay(x):

zig case: if x is child of root do left rotation or right rotation around parent

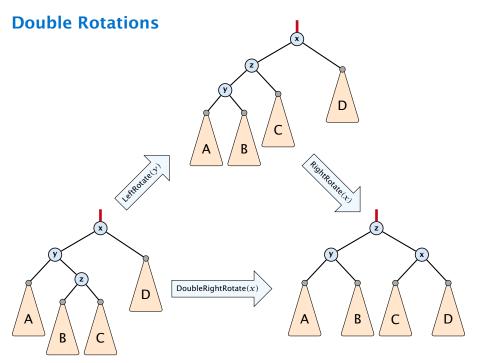
Splay: Zigzag Case



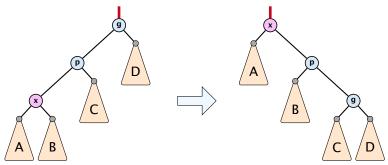
better option splay(x):

- zigzag case: if x is right child and parent of x is left child (or x left child parent of x right child)
- do double right rotation around grand-parent (resp. double left rotation)

7.3 Splay Trees 6. Feb. 202



Splay: Zigzig Case

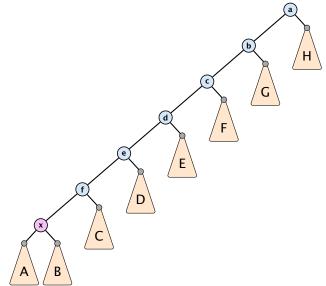


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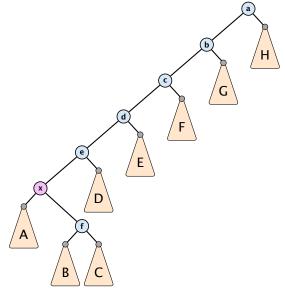
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7.3 Splay Trees

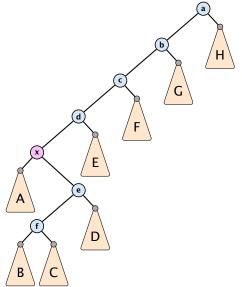
Splay vs. Move to Root

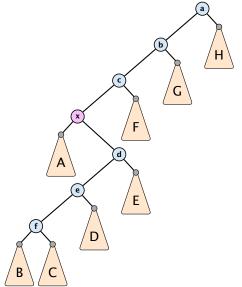


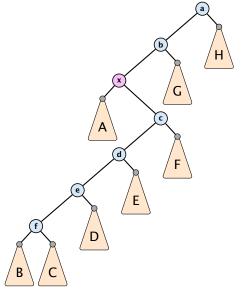
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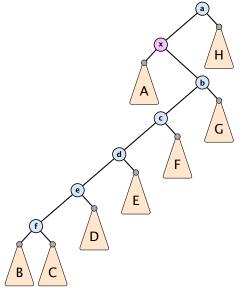
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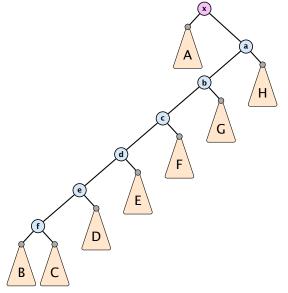


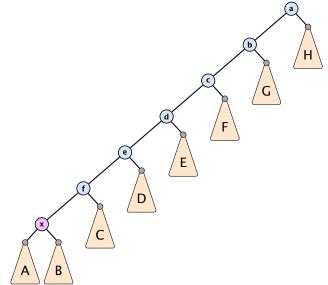


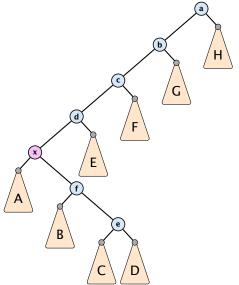


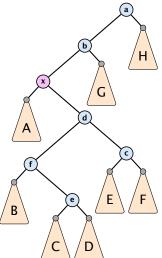


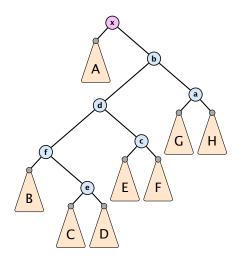












Static Optimality

Suppose we have a sequence of m find-operations. find(x) appears h_x times in this sequence.

The cost of a static search tree *T* is:

$$cost(T) = m + \sum_{x} h_{x} \operatorname{depth}_{T}(x)$$

The total cost for processing the sequence on a splay-tree is $\mathcal{O}(\cos(T_{\min}))$, where T_{\min} is an optimal static search tree.

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Dynamic Optimality

Let S be a sequence with m find-operations.

Let A be a data-structure based on a search tree:

- the cost for accessing element x is 1 + depth(x);
- after accessing x the tree may be re-arranged through rotations;

Conjecture:

A splay tree that only contains elements from S has cost $\mathcal{O}(\cos t(A,S))$, for processing S.

Lemma 5

Splay Trees have an amortized running time of $O(\log n)$ for all operations.

Amortized Analysis

Definition 6

A data structure with operations $op_1(), \ldots, op_k()$ has amortized running times t_1, \ldots, t_k for these operations if the following holds.

Suppose you are given a sequence of operations (starting with an empty data-structure) that operate on at most n elements, and let k_i denote the number of occurences of $\operatorname{op}_i()$ within this sequence. Then the actual running time must be at most $\sum_i k_i \cdot t_i(n)$.

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Then

$$\sum_{i=1}^{k} c_i \le \sum_{i=1}^{k} c_i + \Phi(D_k) - \Phi(D_0) = \sum_{i=1}^{k} \hat{c}_i$$

This means the amortized costs can be used to derive a bound on the total cost.

Stack

- **►** *S.* push()
- ► *S.* pop()
- ► *S.* multipop(*k*): removes *k* items from the stack. If the stack currently contains less than *k* items it empties the stack.
- ► The user has to ensure that pop and multipop do not generate an underflow.

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Actual cost:

- ► *S.* push(): cost 1.
- ► *S.* pop(): cost 1.
- *S.* multipop(k): cost min{size, k} = k.

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► **S. pop()**: cost

$$\hat{C}_{\mathrm{pop}} = C_{\mathrm{pop}} + \Delta \Phi = 1 - 1 \leq 0 \ .$$

 \triangleright S. multipop(k): cost

$$\hat{C}_{mn} = C_{mn} + \Delta \Phi = \min\{\text{size}, k\} - \min\{\text{size}, k\} \le 0$$
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Incrementing a binary counter:

Consider a computational model where each bit-operation costs one time-unit.

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Actual cost:

- ► Changing bit from 0 to 1: cost 1.
- Changing bit from 1 to 0: cost 1.
- ▶ Increment: cost is k + 1, where k is the number of consecutive ones in the least significant bit-positions (e.g, 001101 has k = 1).



Choose potential function $\Phi(x)=k$, where k denotes the number of ones in the binary representation of x.

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$$\hat{C}_{0\to 1} = C_{0\to 1} + \Delta \Phi = 1 + 1 \le 2 .$$

► Changing bit from 1 to 0:

$$\hat{C}_{1\to 0} = C_{1\to 0} + \Delta \Phi = 1 - 1 \le 0 \ .$$

Choose potential function $\Phi(x) = k$, where k denotes the number of ones in the binary representation of x.

Amortized cost:

► Changing bit from 0 to 1:

$$\hat{C}_{0\to 1} = C_{0\to 1} + \Delta \Phi = 1 + 1 \le 2 .$$

► Changing bit from 1 to 0:

$$\hat{C}_{1\to 0} = C_{1\to 0} + \Delta \Phi = 1 - 1 \le 0$$
.

▶ Increment: Let k denotes the number of consecutive ones in the least significant bit-positions. An increment involves k (1 \rightarrow 0)-operations, and one (0 \rightarrow 1)-operation.

Hence, the amortized cost is $k\hat{C}_{1\rightarrow 0} + \hat{C}_{0\rightarrow 1} \leq 2$.

Splay Trees

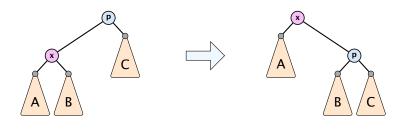
potential function for splay trees:

- ightharpoonup size $s(x) = |T_x|$
- $rank r(x) = \log_2(s(x))$

amortized cost = real cost + potential change

The cost is essentially the cost of the splay-operation, which is 1 plus the number of rotations.

Splay: Zig Case



 $\Delta\Phi =$

Splay: Zig Case



$$\Delta\Phi = r'(x) + r'(p) - r(x) - r(p)$$



$$\Delta\Phi = r'(x) + r'(p) - r(x) - r(p)$$
$$= r'(p) - r(x)$$

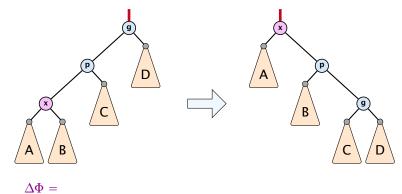


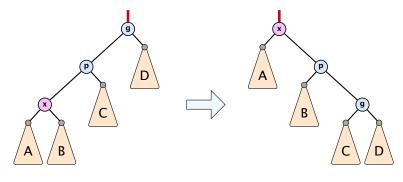
$$\Delta \Phi = r'(x) + r'(p) - r(x) - r(p)$$
$$= r'(p) - r(x)$$
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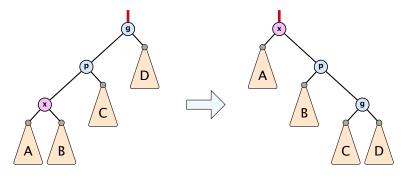
$$\Delta\Phi = r'(x) + r'(p) - r(x) - r(p)$$
$$= r'(p) - r(x)$$
$$\leq r'(x) - r(x)$$

$$cost_{ziq} \le 1 + 3(r'(x) - r(x))$$



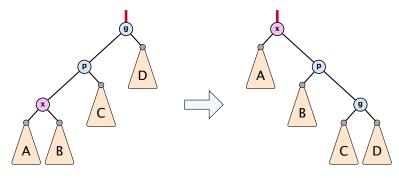


$$\Delta\Phi=r'(x)+r'(p)+r'(g)-r(x)-r(p)-r(g)$$



$$\Delta \Phi = r'(x) + r'(p) + r'(g) - r(x) - r(p) - r(g)$$

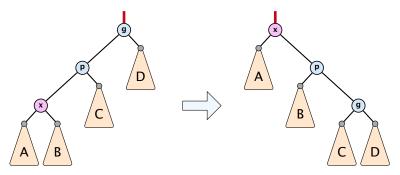
= $r'(p) + r'(g) - r(x) - r(p)$



$$\Delta \Phi = r'(x) + r'(p) + r'(g) - r(x) - r(p) - r(g)$$

$$= r'(p) + r'(g) - r(x) - r(p)$$

$$\leq r'(x) + r'(g) - r(x) - r(x)$$

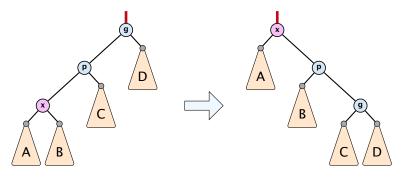


$$\Delta\Phi = r'(x) + r'(p) + r'(g) - r(x) - r(p) - r(g)$$

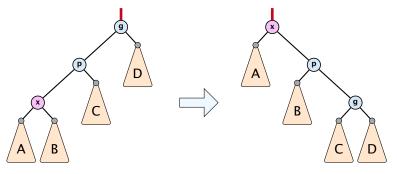
$$= r'(p) + r'(g) - r(x) - r(p)$$

$$\leq r'(x) + r'(g) - r(x) - r(x)$$

$$= r'(x) + r'(g) + r(x) - 3r'(x) + 3r'(x) - r(x) - 2r(x)$$



$$\begin{split} \Delta \Phi &= r'(x) + r'(p) + r'(g) - r(x) - r(p) - r(g) \\ &= r'(p) + r'(g) - r(x) - r(p) \\ &\leq r'(x) + r'(g) - r(x) - r(x) \\ &= r'(x) + r'(g) + r(x) - 3r'(x) + 3r'(x) - r(x) - 2r(x) \\ &= -2r'(x) + r'(g) + r(x) + 3(r'(x) - r(x)) \end{split}$$



$$\Delta\Phi = r'(x) + r'(p) + r'(g) - r(x) - r(p) - r(g)$$

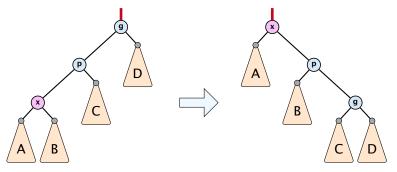
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$$= -2r'(x) + r'(g) + r(x) + 3(r'(x) - r(x))$$

$$\leq -2 + 3(r'(x) - r(x))$$



$$\Delta\Phi = r'(x) + r'(p) + r'(g) - r(x) - r(p) - r(g)$$

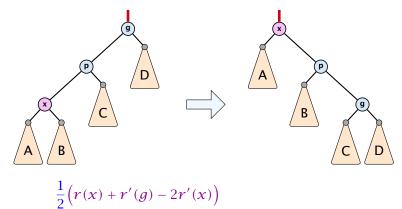
$$= r'(p) + r'(g) - r(x) - r(p)$$

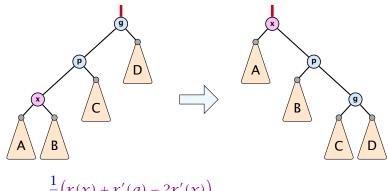
$$\leq r'(x) + r'(g) - r(x) - r(x)$$

$$= r'(x) + r'(g) + r(x) - 3r'(x) + 3r'(x) - r(x) - 2r(x)$$

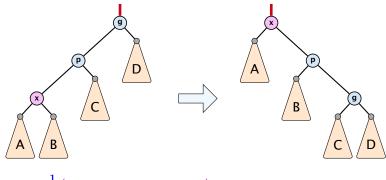
$$= -2r'(x) + r'(g) + r(x) + 3(r'(x) - r(x))$$

$$\leq -2 + 3(r'(x) - r(x)) \Rightarrow \cos t_{zigzig} \leq 3(r'(x) - r(x))$$





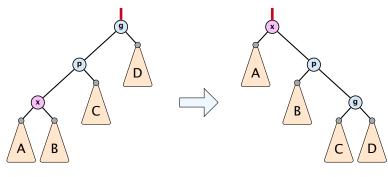
$$\frac{1}{2} \Big(r(x) + r'(g) - 2r'(x) \Big) \\
= \frac{1}{2} \Big(\log(s(x)) + \log(s'(g)) - 2\log(s'(x)) \Big)$$



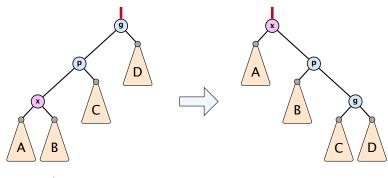
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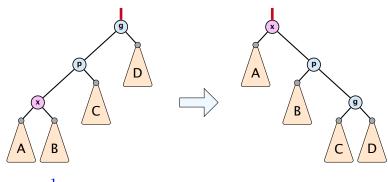
$$= \frac{1}{2} \log \left(\frac{s(x)}{s'(x)} \right) + \frac{1}{2} \log \left(\frac{s'(g)}{s'(x)} \right)$$



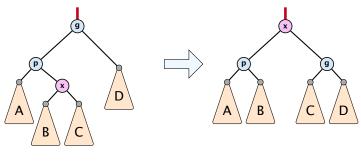
$$\begin{split} \frac{1}{2} \Big(r(x) + r'(g) - 2r'(x) \Big) \\ &= \frac{1}{2} \Big(\log(s(x)) + \log(s'(g)) - 2\log(s'(x)) \Big) \\ &= \frac{1}{2} \log \Big(\frac{s(x)}{s'(x)} \Big) + \frac{1}{2} \log \Big(\frac{s'(g)}{s'(x)} \Big) \\ &\leq \log \Big(\frac{1}{2} \frac{s(x)}{s'(x)} + \frac{1}{2} \frac{s'(g)}{s'(x)} \Big) \end{split}$$



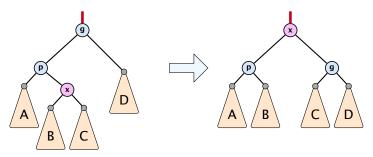
$$\begin{split} &\frac{1}{2}\Big(r(x)+r'(g)-2r'(x)\Big)\\ &=\frac{1}{2}\Big(\log(s(x))+\log(s'(g))-2\log(s'(x))\Big)\\ &=\frac{1}{2}\log\Big(\frac{s(x)}{s'(x)}\Big)+\frac{1}{2}\log\Big(\frac{s'(g)}{s'(x)}\Big)\\ &\leq\log\Big(\frac{1}{2}\frac{s(x)}{s'(x)}+\frac{1}{2}\frac{s'(g)}{s'(x)}\Big)\leq\log\Big(\frac{1}{2}\Big) \end{split}$$



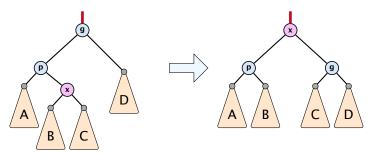
$$\begin{split} \frac{1}{2} \Big(r(x) + r'(g) - 2r'(x) \Big) \\ &= \frac{1}{2} \Big(\log(s(x)) + \log(s'(g)) - 2\log(s'(x)) \Big) \\ &= \frac{1}{2} \log \Big(\frac{s(x)}{s'(x)} \Big) + \frac{1}{2} \log \Big(\frac{s'(g)}{s'(x)} \Big) \\ &\leq \log \Big(\frac{1}{2} \frac{s(x)}{s'(x)} + \frac{1}{2} \frac{s'(g)}{s'(x)} \Big) \leq \log \Big(\frac{1}{2} \Big) = -1 \end{split}$$





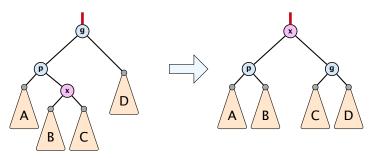


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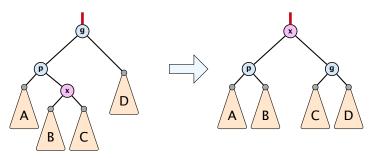
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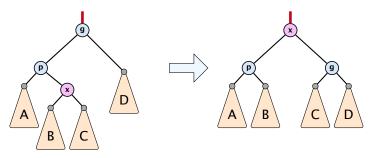


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$$\leq r'(p) + r'(g) - r(x) - r(x)$$

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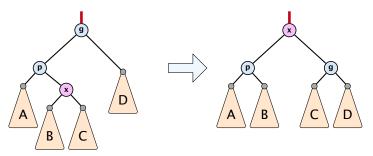
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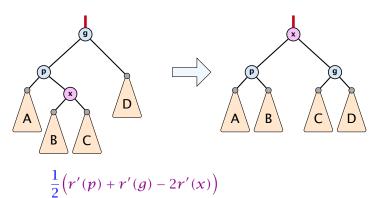
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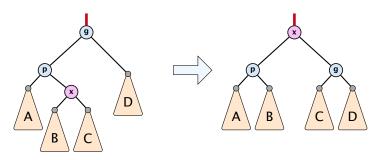
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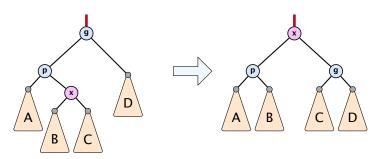
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$$\leq -2 + 2(r'(x) - r(x)) \Rightarrow cost_{zigzag} \leq 3(r'(x) - r(x))$$

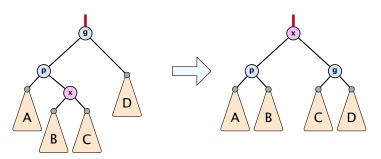




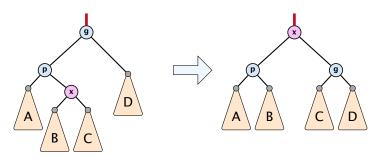
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Amortized cost of the whole splay operation:

$$\leq 1 + 1 + \sum_{\text{steps } t} 3(r_t(x) - r_{t-1}(x))$$

$$= 2 + 3(r(\text{root}) - r_0(x))$$

$$\leq \mathcal{O}(\log n)$$

Suppose you want to develop a data structure with:

- Insert(x): insert element x.
- **Search**(k): search for element with key k.
- **Delete**(x): delete element referenced by pointer x.
- ▶ find-by-rank(ℓ): return the ℓ -th element; return "error" if the data-structure contains less than ℓ elements.

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Augment an existing data-structure instead of developing a new one.

How to augment a data-structure

1. choose an underlying data-structure

- Of course, the above steps heavily depend on each other. For example it makes no sense to choose additional information to be stored (Step 2), and later realize that either the information cannot be maintained efficiently (Step 3) or is not sufficient to support the new operations (Step 4).
- However, the above outline is a good way to describe/document a new data-structure.

How to augment a data-structure

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- determine additional information to be stored in the underlying structure

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How to augment a data-structure

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- 4. develop the new operations
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- However, the above outline is a good way to describe/document a new data-structure.

Goal: Design a data-structure that supports insert, delete, search, and find-by-rank in time $O(\log n)$.

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- 1. We choose a red-black tree as the underlying data-structure.
- 2. We store in each node v the size of the sub-tree rooted at v.
- 3. We need to be able to update the size-field in each node without asymptotically affecting the running time of insert, delete, and search. We come back to this step later...

Goal: Design a data-structure that supports insert, delete, search, and find-by-rank in time $O(\log n)$.

4. How does find-by-rank work? Find-by-rank(k) = Select(root,k) with

```
Algorithm 1 Select(x, i)
```

```
1: if x = \text{null} then return error
```

2: **if** $left[x] \neq null$ **then** $r \leftarrow left[x]$. size + 1 **else** $r \leftarrow 1$

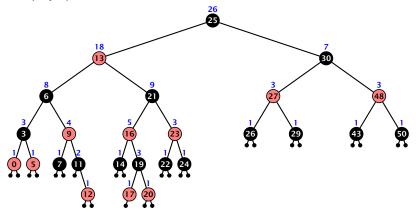
3: **if** i = r **then return** x

4: if i < r then

5: **return** Select(left[x], i)

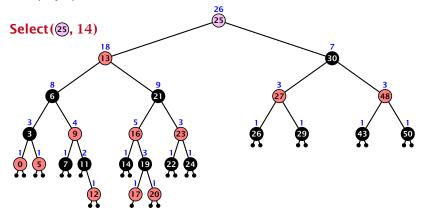
6: **else**

7: **return** Select(right[x], i - r)



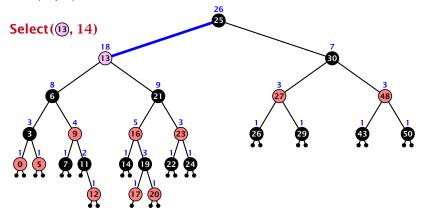
- decide whether you have to proceed into the left or right sub-tree
- adjust the rank that you are searching for if you go right





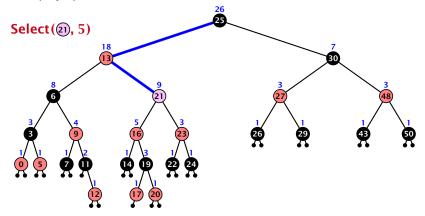
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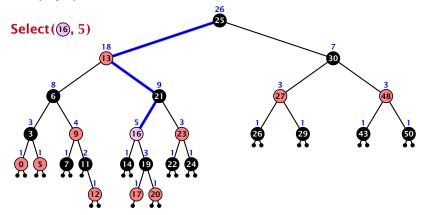
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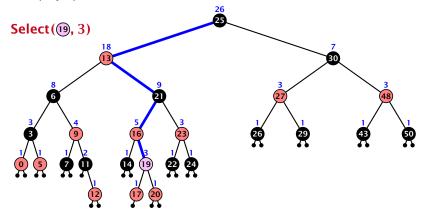
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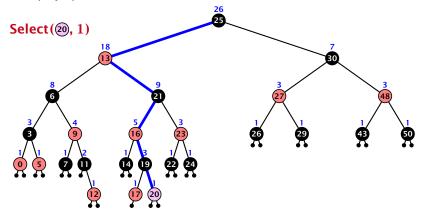
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Goal: Design a data-structure that supports insert, delete, search, and find-by-rank in time $\mathcal{O}(\log n)$.

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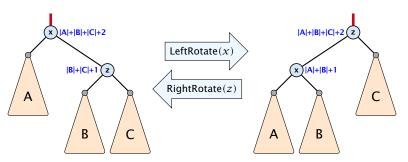
Search(k): Nothing to do.

Insert(x): When going down the search path increase the size field for each visited node. Maintain the size field during rotations.

Delete(*x*): Directly after splicing out a node traverse the path from the spliced out node upwards, and decrease the size counter on every node on this path. Maintain the size field during rotations.

Rotations

The only operation during the fix-up procedure that alters the tree and requires an update of the size-field:



The nodes x and z are the only nodes changing their size-fields.

The new size-fields can be computed locally from the size-fields of the children.

Why do we not use a list for implementing the ADT Dynamic Set?

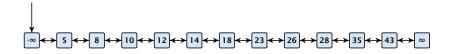
Why do we not use a list for implementing the ADT Dynamic Set?

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6. Feb. 2022

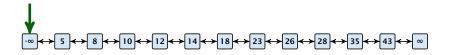
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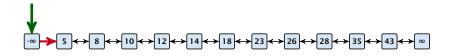
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6. Feb. 2022

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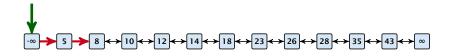
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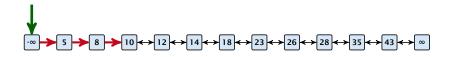
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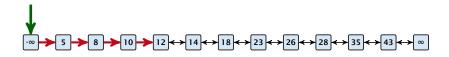
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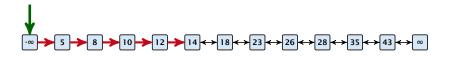
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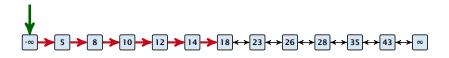
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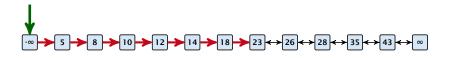
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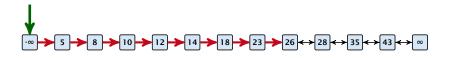
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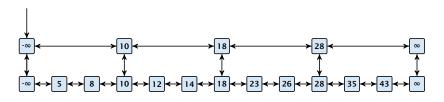
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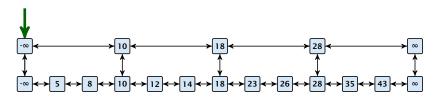
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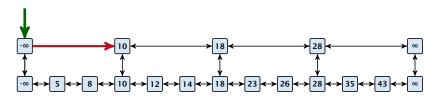
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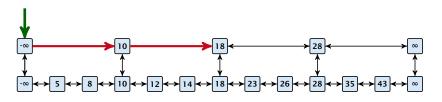
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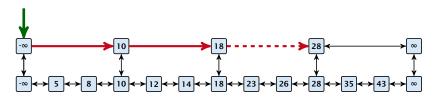
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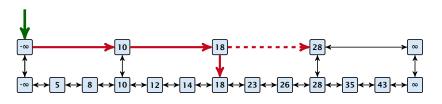


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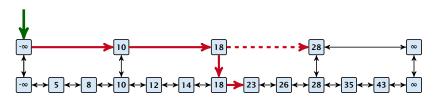
How can we improve the search-operation?

Add an express lane:



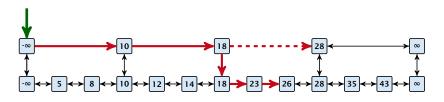
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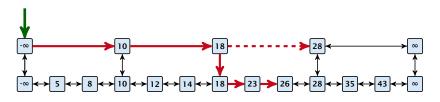
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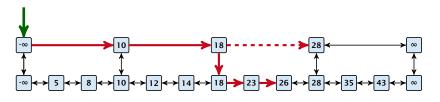
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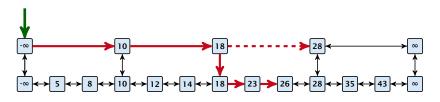


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Choose $|L_1| = \sqrt{n}$. Then search time $\Theta(\sqrt{n})$.

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Choosing $k = \Theta(\log n)$ gives a logarithmic running time.

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Use randomization instead!

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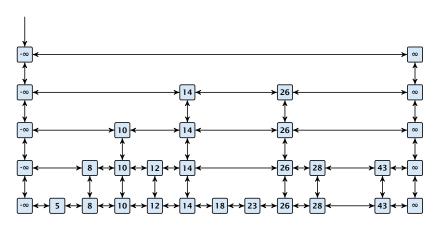
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The time for both operations is dominated by the search time.

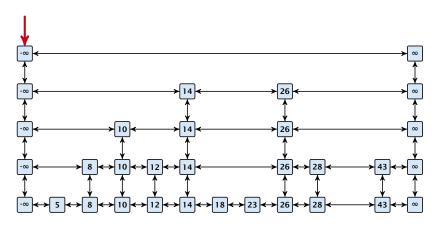
Insert (35):





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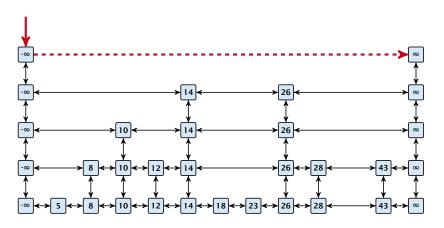
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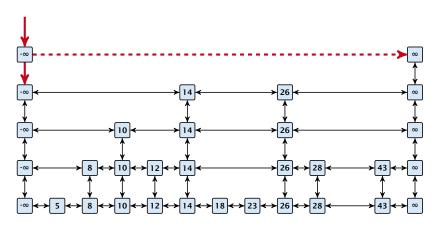
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Insert (35):



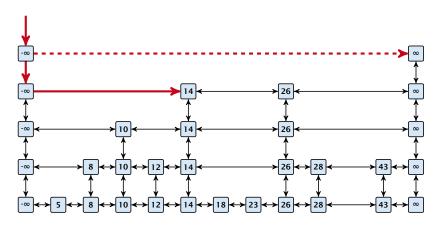


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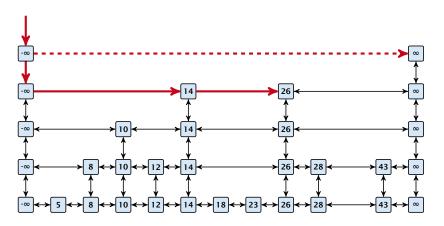


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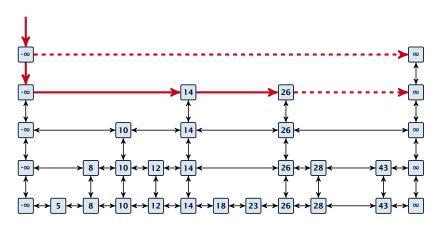


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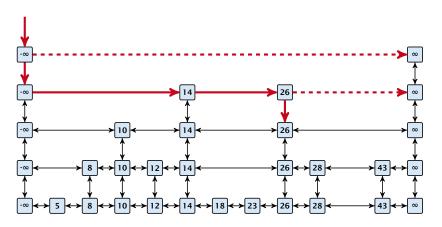


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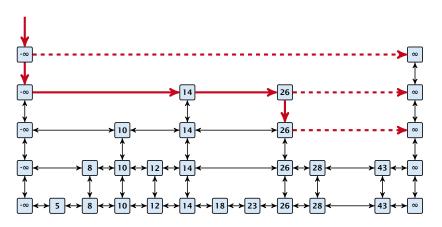
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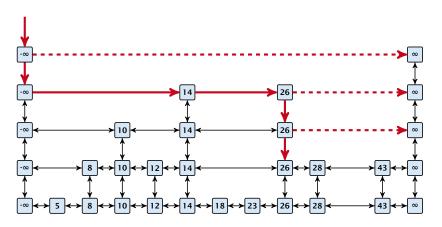


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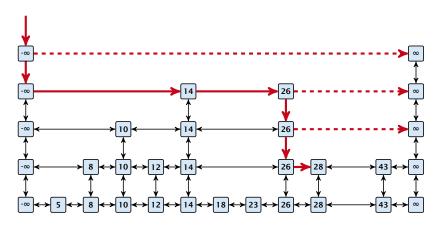


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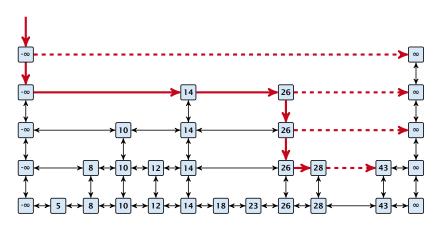


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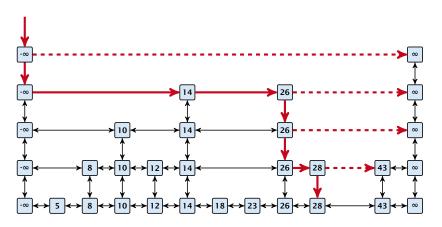


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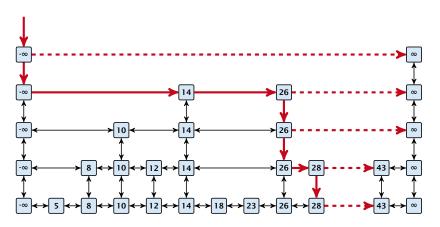
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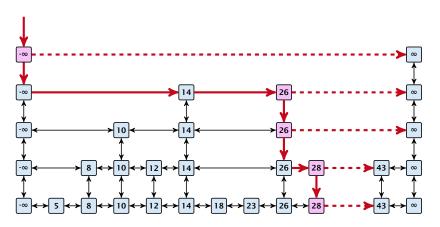
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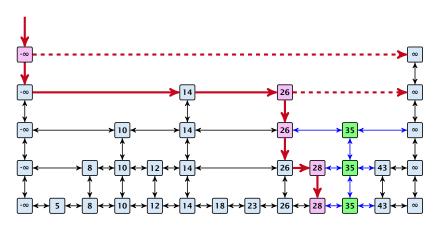


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Definition 7 (High Probability)

We say a **randomized** algorithm has running time $\mathcal{O}(\log n)$ with high probability if for any constant α the running time is at most $\mathcal{O}(\log n)$ with probability at least $1 - \frac{1}{n^{\alpha}}$.

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Here the \mathcal{O} -notation hides a constant that may depend on α .



Suppose there are polynomially many events E_1, E_2, \dots, E_ℓ , $\ell = n^c$ each holding with high probability (e.g. E_i may be the event that the i-th search in a skip list takes time at most $\mathcal{O}(\log n)$).

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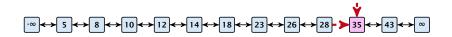
This means $\Pr[E_1 \wedge \cdots \wedge E_{\ell}]$ holds with high probability.

Lemma 8

A search (and, hence, also insert and delete) in a skip list with n elements takes time O(logn) with high probability (w. h. p.).

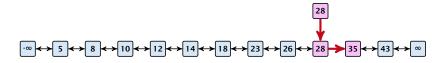
$$\begin{array}{c} -\infty \longleftrightarrow 5 \longleftrightarrow 8 \longleftrightarrow 10 \longleftrightarrow 12 \longleftrightarrow 14 \longleftrightarrow 18 \longleftrightarrow 23 \longleftrightarrow 26 \longleftrightarrow 28 \longleftrightarrow 35 \longleftrightarrow 43 \longleftrightarrow \infty \end{array}$$

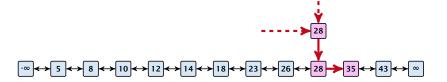
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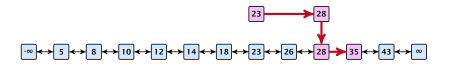


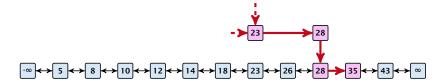
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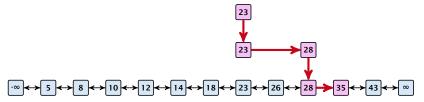


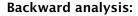


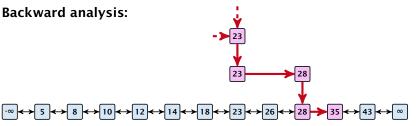






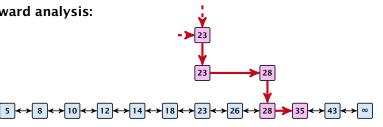




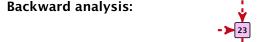


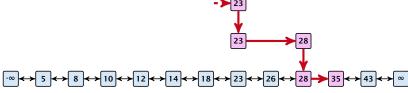


Backward analysis:



At each point the path goes up with probability 1/2 and left with probability 1/2.

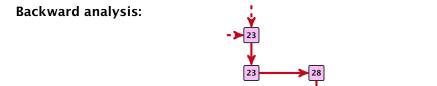




At each point the path goes up with probability 1/2 and left with probability 1/2.

We show that w.h.p:

A "long" search path must also go very high.

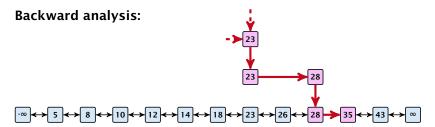


At each point the path goes up with probability 1/2 and left with probability 1/2.

 \leftrightarrow 8 \leftrightarrow 10 \leftrightarrow 12 \leftrightarrow 14 \leftrightarrow 18 \leftrightarrow 23 \leftrightarrow 26 \leftrightarrow 28

We show that w.h.p:

- A "long" search path must also go very high.
- There are no elements in high lists.



At each point the path goes up with probability 1/2 and left with probability 1/2.

We show that w.h.p:

- A "long" search path must also go very high.
- There are no elements in high lists.

From this it follows that w.h.p. there are no long paths.



Estimation for Binomial Coefficients

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

$$\binom{n}{k}$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

$$\binom{n}{k} = \frac{n!}{k! \cdot (n-k)!}$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

$$\binom{n}{k} = \frac{n!}{k! \cdot (n-k)!} = \frac{n \cdot \ldots \cdot (n-k+1)}{k \cdot \ldots \cdot 1}$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

$$\binom{n}{k} = \frac{n!}{k! \cdot (n-k)!} = \frac{n \cdot \ldots \cdot (n-k+1)}{k \cdot \ldots \cdot 1} \ge \left(\frac{n}{k}\right)^k$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

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$$\binom{n}{k} = \frac{n \cdot \ldots \cdot (n-k+1)}{k!}$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

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$$\binom{n}{k} = \frac{n \cdot \ldots \cdot (n-k+1)}{k!} \le \frac{n^k}{k!}$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

$$\binom{n}{k} = \frac{n!}{k! \cdot (n-k)!} = \frac{n \cdot \ldots \cdot (n-k+1)}{k \cdot \ldots \cdot 1} \ge \left(\frac{n}{k}\right)^k$$

$$\binom{n}{k} = \frac{n \cdot \ldots \cdot (n-k+1)}{k!} \le \frac{n^k}{k!} = \frac{n^k \cdot k^k}{k^k \cdot k!}$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

$$\binom{n}{k} = \frac{n!}{k! \cdot (n-k)!} = \frac{n \cdot \ldots \cdot (n-k+1)}{k \cdot \ldots \cdot 1} \ge \left(\frac{n}{k}\right)^k$$

$$\binom{n}{k} = \frac{n \cdot \dots \cdot (n-k+1)}{k!} \le \frac{n^k}{k!} = \frac{n^k \cdot k^k}{k^k \cdot k!}$$
$$= \left(\frac{n}{k}\right)^k \cdot \frac{k^k}{k!}$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

$$\binom{n}{k} = \frac{n!}{k! \cdot (n-k)!} = \frac{n \cdot \ldots \cdot (n-k+1)}{k \cdot \ldots \cdot 1} \ge \left(\frac{n}{k}\right)^k$$

$$\binom{n}{k} = \frac{n \cdot \dots \cdot (n - k + 1)}{k!} \le \frac{n^k}{k!} = \frac{n^k \cdot k^k}{k^k \cdot k!}$$
$$= \left(\frac{n}{k}\right)^k \cdot \frac{k^k}{k!} \le \left(\frac{n}{k}\right)^k \cdot \sum_{i \ge 0} \frac{k^i}{i!}$$

$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

$$\binom{n}{k} = \frac{n!}{k! \cdot (n-k)!} = \frac{n \cdot \ldots \cdot (n-k+1)}{k \cdot \ldots \cdot 1} \ge \left(\frac{n}{k}\right)^k$$

$$\binom{n}{k} = \frac{n \cdot \dots \cdot (n-k+1)}{k!} \le \frac{n^k}{k!} = \frac{n^k \cdot k^k}{k^k \cdot k!}$$
$$= \left(\frac{n}{k}\right)^k \cdot \frac{k^k}{k!} \le \left(\frac{n}{k}\right)^k \cdot \sum_{i=0}^k \frac{k^i}{i!} = \left(\frac{en}{k}\right)^k$$

Let $E_{z,k}$ denote the event that a search path is of length z (number of edges) but does not visit a list above L_k .

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In particular, this means that during the construction in the backward analysis we see at most k heads (i.e., coin flips that tell you to go up) in z trials.

 $\Pr[E_{z,k}]$

 $Pr[E_{z,k}] \leq Pr[at most k heads in z trials]$

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$$\leq \binom{z}{k} 2^{-(z-k)}$$

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$$\leq \binom{z}{k} 2^{-(z-k)} \leq \left(\frac{ez}{k}\right)^k 2^{-(z-k)}$$

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choosing $k = y \log n$ with $y \ge 1$ and $z = (\beta + \alpha)y \log n$

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$$\leq \left(\frac{2ez}{k}\right)^k 2^{-\beta k} \cdot n^{-y\alpha} \leq \left(\frac{2ez}{2^\beta k}\right)^k \cdot n^{-\alpha}$$

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$$\leq \binom{z}{k} 2^{-(z-k)} \leq \left(\frac{ez}{k}\right)^k 2^{-(z-k)} \leq \left(\frac{2ez}{k}\right)^k 2^{-z}$$
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now choosing $\beta = 6\alpha$ gives

 $\Pr[E_{z,k}] \leq \Pr[\text{at most } k \text{ heads in } z \text{ trials}]$

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$$\leq \left(\frac{2e(\beta + \alpha)}{2^\beta}\right)^k n^{-\alpha}$$
 now choosing $\beta = 6\alpha$ gives

$$(42\alpha)^k$$

$$\leq \left(\frac{42\alpha}{64^{\alpha}}\right)^k n^{-\alpha}$$

 $Pr[E_{z,k}] \leq Pr[at most k heads in z trials]$

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 now choosing $\beta = 6\alpha$ gives

$$\leq \left(\frac{42\alpha}{64^{\alpha}}\right)^k n^{-\alpha} \leq n^{-\alpha}$$

 $Pr[E_{z,k}] \leq Pr[at most k heads in z trials]$

$$\leq \binom{z}{k} 2^{-(z-k)} \leq \left(\frac{ez}{k}\right)^k 2^{-(z-k)} \leq \left(\frac{2ez}{k}\right)^k 2^{-z}$$
 choosing $k = y \log n$ with $y \geq 1$ and $z = (\beta + \alpha) y \log n$
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now choosing $\beta = 6\alpha$ gives

$$\leq \left(\frac{42\alpha}{64\alpha}\right)^k n^{-\alpha} \leq n^{-\alpha}$$

for $\alpha \geq 1$.

So far we fixed $k = y \log n$, $y \ge 1$, and $z = 7\alpha y \log n$, $\alpha \ge 1$.

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This means that a search path of length $\Omega(\log n)$ visits a list on a level $\Omega(\log n)$, w.h.p.

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Let A_{k+1} denote the event that the list L_{k+1} is non-empty. Then

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Let A_{k+1} denote the event that the list L_{k+1} is non-empty. Then

$$\Pr[A_{k+1}] \le n2^{-(k+1)} \le n^{-(\gamma-1)}$$
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For the search to take at least $z = 7\alpha y \log n$ steps either the event $E_{z,k}$ or the event A_{k+1} must hold.

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For the search to take at least $z = 7\alpha\gamma \log n$ steps either the event $E_{z,k}$ or the event A_{k+1} must hold. Hence,

Pr[search requires z steps]

So far we fixed $k = y \log n$, $y \ge 1$, and $z = 7\alpha y \log n$, $\alpha \ge 1$.

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 $\Pr[\text{search requires } z \text{ steps}] \leq \Pr[E_{z,k}] + \Pr[A_{k+1}]$

So far we fixed $k = y \log n$, $y \ge 1$, and $z = 7\alpha y \log n$, $\alpha \ge 1$.

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$$\Pr[\text{search requires } z \text{ steps}] \le \Pr[E_{z,k}] + \Pr[A_{k+1}]$$

 $\le n^{-\alpha} + n^{-(\gamma-1)}$

7.5 Skip Lists

So far we fixed $k = y \log n$, $y \ge 1$, and $z = 7\alpha y \log n$, $\alpha \ge 1$.

This means that a search path of length $\Omega(\log n)$ visits a list on a level $\Omega(\log n)$, w.h.p.

Let A_{k+1} denote the event that the list L_{k+1} is non-empty. Then

$$\Pr[A_{k+1}] \le n2^{-(k+1)} \le n^{-(\gamma-1)}$$
.

For the search to take at least $z = 7\alpha\gamma \log n$ steps either the event $E_{z,k}$ or the event A_{k+1} must hold. Hence,

$$\Pr[\text{search requires } z \text{ steps}] \le \Pr[E_{z,k}] + \Pr[A_{k+1}]$$

 $\le n^{-\alpha} + n^{-(\gamma-1)}$

This means, the search requires at most z steps, w.h.p.

7.6 van Emde Boas Trees

Dynamic Set Data Structure *S***:**

- \triangleright S. insert(x)
- \triangleright S. delete(x)
- \triangleright S. search(x)
- ► *S*. min()
- ► *S*. max()
- \triangleright S. succ(x)
- \triangleright S. pred(x)

7.6 van Emde Boas Trees

For this chapter we ignore the problem of storing satellite data:

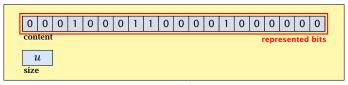
- \triangleright S. insert(x): Inserts x into S.
- ▶ S. delete(x): Deletes x from S. Usually assumes that $x \in S$.
- **S.** member(x): Returns 1 if $x \in S$ and 0 otw.
- **S.** min(): Returns the value of the minimum element in S.
- **S.** $\max()$: Returns the value of the maximum element in S.
- ► *S.* succ(*x*): Returns successor of *x* in *S*. Returns null if *x* is maximum or larger than any element in *S*. Note that *x* needs not to be in *S*.
- ▶ S. pred(x): Returns the predecessor of x in S. Returns null if x is minimum or smaller than any element in S. Note that x needs not to be in S.



7.6 van Emde Boas Trees

Can we improve the existing algorithms when the keys are from a restricted set?

In the following we assume that the keys are from $\{0, 1, \dots, u-1\}$, where u denotes the size of the universe.



one array of *u* bits

Use an array that encodes the indicator function of the dynamic set.

```
Algorithm 1 array.insert(x)
```

1: content[x] \leftarrow 1;

Algorithm 2 array.delete(x)

1: content[x] \leftarrow 0;

Algorithm 3 array.member(x)

1: **return** content[x];

- Note that we assume that x is valid, i.e., it falls within the array boundaries.
- Obviously(?) the running time is constant.

Algorithm 4 array.max()

1: for $(i = \text{size} -1; i \ge 0; i--)$ do 2: if content[i] = 1 then return i;

3: return null;

Algorithm 4 array.max()

1: for $(i = \text{size} - 1; i \ge 0; i--)$ do 2: if content[i] = 1 then return i;

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Algorithm 5 array.min()

```
1: for (i = 0; i < \text{size}; i++) do
```

2: **if** content[i] = 1 **then return** i;

3: return null;

Algorithm 4 array.max()

```
1: for (i = \text{size} -1; i \ge 0; i--) do
2: if content[i] = 1 then return i;
```

3: return null;

Algorithm 5 array.min()

1: **for** (i = 0; i < size; i++) **do** 2: **if** content[i] = 1 **then return** i;

3: return null:

Running time is $\mathcal{O}(u)$ in the worst case.

Algorithm 6 array.succ(x)

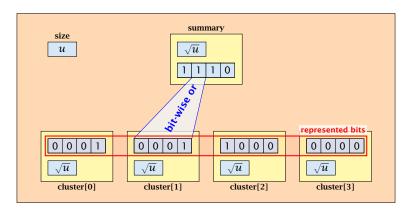
1: for (i = x + 1; i < size; i++) do 2: if content[i] = 1 then return i; 3: return null;

Algorithm 7 array.pred(x)

1: for $(i = x - 1; i \ge 0; i--)$ do 2: if content[i] = 1 then return i;

3: return null:

Running time is $\mathcal{O}(u)$ in the worst case.



- \sqrt{u} cluster-arrays of \sqrt{u} bits.
- One summary-array of \sqrt{u} bits. The *i*-th bit in the summary array stores the bit-wise or of the bits in the *i*-th cluster.

The bit for a key x is contained in cluster number $\left\lfloor \frac{x}{\sqrt{u}} \right\rfloor$.

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Within the cluster-array the bit is at position $x \mod \sqrt{u}$.

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Within the cluster-array the bit is at position $x \mod \sqrt{u}$.

For simplicity we assume that $u=2^{2k}$ for some $k\geq 1$. Then we can compute the cluster-number for an entry x as $\mathrm{high}(x)$ (the upper half of the dual representation of x) and the position of x within its cluster as $\mathrm{low}(x)$ (the lower half of the dual representation).

Algorithm 8 member(x)

1: **return** cluster[high(x)]. member(low(x));

Algorithm 8 member(x)

1: **return** cluster[high(x)]. member(low(x));

Algorithm 9 insert(x)

- 1: $\operatorname{cluster}[\operatorname{high}(x)].\operatorname{insert}(\operatorname{low}(x));$
- 2: summary.insert(high(x));

Algorithm 8 member(x)

1: **return** cluster[high(x)]. member(low(x));

Algorithm 9 insert(x)

- 1: cluster[high(x)].insert(low(x));
- 2: $\operatorname{summary.insert}(\operatorname{high}(x));$
- ► The running times are constant, because the corresponding array-functions have constant running times.

Algorithm 10 delete(x)

- 1: cluster[high(x)].delete(low(x));
- 2: **if** cluster[high(x)]. min() = null **then**
- 3: summary . delete(high(x));

Algorithm 10 delete(x)

- 1: cluster[high(x)]. delete(low(x));
- 2: **if** cluster[high(x)].min() = null **then**
- 3: summary . delete(high(x));
- The running time is dominated by the cost of a minimum computation on an array of size \sqrt{u} . Hence, $\mathcal{O}(\sqrt{u})$.

Algorithm 11 max()

- 1: maxcluster ← summary.max(); 2: if maxcluster = null return null; 3: offs ← cluster[maxcluster].max() 4: return maxcluster ∘ offs;

Algorithm 11 max()

- 1: *maxcluster* ← summary.max();
- 2: **if** *maxcluster* = null **return** null;
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Algorithm 12 min()

- 1: *mincluster* ← summary.min();
- 2: **if** *mincluster* = null **return** null;
- 3: $offs \leftarrow cluster[mincluster].min();$
- 4: **return** *mincluster* ∘ *offs*;

Algorithm 11 max()

- 1: *maxcluster* ← summary.max();
- 2: **if** *maxcluster* = null **return** null;
- 3: offs ← cluster[maxcluster].max()
- 4: **return** *maxcluster* ∘ *offs*;

Algorithm 12 min()

- 1: *mincluster* ← summary.min();
- 2: **if** *mincluster* = null **return** null;
- 3: $offs \leftarrow cluster[mincluster].min();$
- 4: **return** *mincluster* ∘ *offs*;

Running time is roughly $2\sqrt{u} = \mathcal{O}(\sqrt{u})$ in the worst case.

The operator \circ stands for the concatenation of two bitstrings.

This means if $x = 0111_2$ and $y = 0001_2$ then $x \circ y = 01110001_2$.

```
Algorithm 13 succ(x)

1: m \leftarrow cluster[high(x)].succ(low(x))

2: if m \neq null then return high(x) \circ m;

3: succcluster \leftarrow summary.succ(high(x));

4: if succcluster \neq null then

5: offs \leftarrow cluster[succcluster].min();

6: return succcluster \circ offs;

7: return null;
```

```
Algorithm 13 \operatorname{succ}(x)

1: m \leftarrow \operatorname{cluster}[\operatorname{high}(x)].\operatorname{succ}(\operatorname{low}(x))

2: if m \neq \operatorname{null} then return \operatorname{high}(x) \circ m;

3: \operatorname{succcluster} \leftarrow \operatorname{summary}.\operatorname{succ}(\operatorname{high}(x));

4: if \operatorname{succcluster} \neq \operatorname{null} then

5: \operatorname{offs} \leftarrow \operatorname{cluster}[\operatorname{succcluster}].\operatorname{min}();

6: \operatorname{return} \operatorname{succcluster} \circ \operatorname{offs};

7: \operatorname{return} \operatorname{null};
```

▶ Running time is roughly $3\sqrt{u} = \mathcal{O}(\sqrt{u})$ in the worst case.

```
Algorithm 14 pred(x)

1: m ← cluster[high(x)].pred(low(x))

2: if m ≠ null then return high(x) ∘ m;

3: predcluster ← summary.pred(high(x));

4: if predcluster ≠ null then

5: offs ← cluster[predcluster].max();

6: return predcluster ∘ offs;

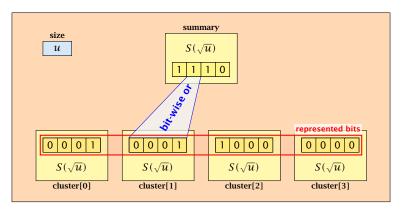
7: return null;
```

• Running time is roughly $3\sqrt{u} = \mathcal{O}(\sqrt{u})$ in the worst case.

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Instead of using sub-arrays, we build a recursive data-structure.

S(u) is a dynamic set data-structure representing u bits:



We assume that $u = 2^{2^k}$ for some k.

The data-structure S(2) is defined as an array of 2-bits (end of the recursion).

The code from Implementation 2 can be used unchanged. We only need to redo the analysis of the running time.

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Note that in the code we do not need to specifically address the non-recursive case. This is achieved by the fact that an S(4) will contain S(2)'s as sub-datastructures, which are arrays. Hence, a call like cluster[1]. min() from within the data-structure S(4) is not a recursive call as it will call the function array. min().

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Note that in the code we do not need to specifically address the non-recursive case. This is achieved by the fact that an S(4) will contain S(2)'s as sub-datastructures, which are arrays. Hence, a call like cluster[1]. $\min()$ from within the data-structure S(4) is not a recursive call as it will call the function $\operatorname{array.min}()$.

This means that the non-recursive case is been dealt with while initializing the data-structure.

Algorithm 15 member(x)

1: **return** cluster[high(x)].member(low(x));

 $T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1.$

Algorithm 16 insert(x)

- 1: $\operatorname{cluster}[\operatorname{high}(x)]$. $\operatorname{insert}(\operatorname{low}(x))$;
- 2: summary.insert(high(x));
- $T_{ins}(u) = 2T_{ins}(\sqrt{u}) + 1.$

Algorithm 17 delete(x)

- 1: $\operatorname{cluster}[\operatorname{high}(x)]$. $\operatorname{delete}(\operatorname{low}(x))$;
- 2: **if** cluster[high(x)].min() = null **then**
- 3: summary . delete(high(x));
- ► $T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1$.

Algorithm 18 min()

- 1: *mincluster* ← summary.min();
- 2: **if** *mincluster* = null **return** null;
- 3: *offs* ← cluster[*mincluster*].min();
- 4: **return** $mincluster \circ offs$;
- $T_{\min}(u) = 2T_{\min}(\sqrt{u}) + 1.$

```
Algorithm 19 succ(x)

1: m \leftarrow cluster[high(x)].succ(low(x))

2: if m \neq null then return high(x) \circ m;

3: succcluster \leftarrow summary.succ(high(x));

4: if succcluster \neq null then

5: offs \leftarrow cluster[succcluster].min();

6: return succcluster \circ offs;

7: return null;
```

 $T_{\text{succ}}(u) = 2T_{\text{succ}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1.$

$$T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$
:

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:

Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{mem}}(2^{\ell})$.

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= $T_{\text{mem}}(2^{\frac{\ell}{2}}) + 1$

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$$X(\ell) = T_{\text{mem}}(2^{\ell}) = T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$

= $T_{\text{mem}}(2^{\frac{\ell}{2}}) + 1 = X(\frac{\ell}{2}) + 1$.

$$T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$
:

Set $\ell := \log u$ and $X(\ell) := T_{\text{mem}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{mem}}(2^{\ell}) = T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$

= $T_{\text{mem}}(2^{\frac{\ell}{2}}) + 1 = X(\frac{\ell}{2}) + 1$.

Using Master theorem gives $X(\ell) = \mathcal{O}(\log \ell)$, and hence $T_{\text{mem}}(u) = \mathcal{O}(\log \log u)$.

$$T_{\rm ins}(u) = 2T_{\rm ins}(\sqrt{u}) + 1.$$

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Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{ins}}(2^{\ell})$. Then

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= $2T_{\text{ins}}(2^{\frac{\ell}{2}}) + 1 = 2X(\frac{\ell}{2}) + 1$.

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Using Master theorem gives $X(\ell) = \mathcal{O}(\ell)$, and hence $T_{\text{ins}}(u) = \mathcal{O}(\log u)$.

The same holds for $T_{\text{max}}(u)$ and $T_{\text{min}}(u)$.

$$T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1 \le 2T_{\text{del}}(\sqrt{u}) + \frac{c}{\log(u)}.$$

$$T_{\rm del}(u) = 2T_{\rm del}(\sqrt{u}) + T_{\rm min}(\sqrt{u}) + 1 \le 2T_{\rm del}(\sqrt{u}) + \frac{c}{\log(u)}.$$

Set $\ell := \log u$ and $X(\ell) := T_{\text{del}}(2^{\ell})$.

$$T_{\rm del}(u) = 2T_{\rm del}(\sqrt{u}) + T_{\rm min}(\sqrt{u}) + 1 \le 2T_{\rm del}(\sqrt{u}) + \frac{c}{\log(u)}.$$

Set $\ell := \log u$ and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

$$T_{\rm del}(u)=2T_{\rm del}(\sqrt{u})+T_{\rm min}(\sqrt{u})+1\leq 2T_{\rm del}(\sqrt{u})+c\log(u).$$
 Set $\ell:=\log u$ and $X(\ell):=T_{\rm del}(2^\ell)$. Then
$$X(\ell)$$

$$T_{\rm del}(u) = 2T_{\rm del}(\sqrt{u}) + T_{\rm min}(\sqrt{u}) + 1 \le 2T_{\rm del}(\sqrt{u}) + \frac{c}{\log(u)}.$$

Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{del}}(2^{\ell})$$

$$T_{\rm del}(u) = 2T_{\rm del}(\sqrt{u}) + T_{\rm min}(\sqrt{u}) + 1 \le 2T_{\rm del}(\sqrt{u}) + \frac{c}{\log(u)}.$$

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$$\ell := \log u$$
 and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

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$$T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1 \le 2T_{\text{del}}(\sqrt{u}) + \frac{c}{\log(u)}.$$

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$$= 2T_{\text{del}}(2^{\frac{\ell}{2}}) + c\ell = 2X(\frac{\ell}{2}) + c\ell .$$

Using Master theorem gives $X(\ell) = \Theta(\ell \log \ell)$, and hence $T_{\text{del}}(u) = \mathcal{O}(\log u \log \log u)$.

$$T_{\rm del}(u) = 2T_{\rm del}(\sqrt{u}) + T_{\rm min}(\sqrt{u}) + 1 \le 2T_{\rm del}(\sqrt{u}) + \frac{c}{\log(u)}.$$

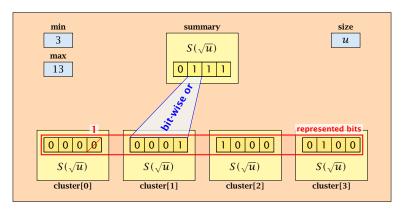
Set $\ell := \log u$ and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{del}}(2^{\ell}) = T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + c \log u$$
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Using Master theorem gives $X(\ell) = \Theta(\ell \log \ell)$, and hence $T_{\text{del}}(u) = \mathcal{O}(\log u \log \log u)$.

The same holds for $T_{\text{pred}}(u)$ and $T_{\text{succ}}(u)$.

Implementation 4: van Emde Boas Trees



- The bit referenced by min is not set within sub-datastructures.
- The bit referenced by max is set within sub-datastructures (if max ≠ min).

Implementation 4: van Emde Boas Trees

Advantages of having max/min pointers:

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Recursive calls for min and max are constant time.

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- Recursive calls for min and max are constant time.
- ightharpoonup min = null means that the data-structure is empty.
- min = max ≠ null means that the data-structure contains exactly one element.
- We can insert into an empty datastructure in constant time by only setting min = max = x.
- We can delete from a data-structure that just contains one element in constant time by setting min = max = null.

Algorithm 20 max()

1: return max;

Algorithm 21 min()

1: return min;

Constant time.

Algorithm 22 member(x)

- 1: **if** $x = \min$ **then return** 1; // TRUE
- 2: **return** cluster[high(x)].member(low(x));
- $T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1 \Longrightarrow T(u) = \mathcal{O}(\log \log u).$

```
Algorithm 23 succ(x)
 1: if min \neq null \wedge x < min then return min:
 2: maxincluster \leftarrow cluster[high(x)].max();
 3: if maxincluster \neq null \land low(x) < maxincluster then
         offs \leftarrow cluster[high(x)]. succ(low(x));
4:
         return high(x) \circ offs;
 5:
6: else
         succeluster \leftarrow summary.succ(high(x));
7:
8:
         if succeluster = null then return null:
9:
         offs \leftarrow cluster[succeluster].min();
         return succeluster o offs:
10:
```

```
Algorithm 35 insert(x)
 1: if min = null then
       \min = x; \max = x;
3: else
4:
       if x < \min then exchange x and \min;
     if x > \max then \max = x;
6:
     if cluster[high(x)]. min = null; then
7:
            summary insert(high(x));
8:
            cluster[high(x)].insert(low(x));
        else
9:
            cluster[high(x)].insert(low(x));
10:
```

 $T_{\text{ins}}(u) = T_{\text{ins}}(\sqrt{u}) + 1 \Longrightarrow T_{\text{ins}}(u) = \mathcal{O}(\log \log u).$

Note that the recusive call in Line 8 takes constant time as the if-condition in Line 6 ensures that we are inserting in an empty sub-tree.

The only non-constant recursive calls are the call in Line 7 and in Line 10. These are mutually exclusive, i.e., only one of these calls will actually occur.

From this we get that $T_{\text{ins}}(u) = T_{\text{ins}}(\sqrt{u}) + 1$.

Assumes that x is contained in the structure.

```
Algorithm 36 delete(x)
1: if min = max then
       min = max = null;
 3: else
4:
       if x = \min then
             firstcluster ← summary.min();
6:
             offs \leftarrow cluster[firstcluster].min();
        x \leftarrow firstcluster \circ offs;
 7:
         \min \leftarrow x:
        cluster[high(x)]. delete(low(x));
 9:
                         continued...
```

Assumes that x is contained in the structure.

```
Algorithm 36 delete(x)
 1: if min = max then
        min = max = null;
 3: else
4:
         if x = \min then
                                              find new minimum
              firstcluster \leftarrow summary.min();
 5:
              offs ← cluster[firstcluster]. min();
6:
              x \leftarrow firstcluster \circ offs;
 7:
 8:
          \min \leftarrow x:
         cluster[high(x)]. delete(low(x));
 9:
                          continued...
```

Assumes that x is contained in the structure.

```
Algorithm 36 delete(x)
 1: if min = max then
        min = max = null;
 3: else
4:
        if x = \min then
              firstcluster \leftarrow summary.min();
 5:
6:
               offs \leftarrow cluster[firstcluster].min();
              x \leftarrow firstcluster \circ offs;
 7:
 8:
              \min \leftarrow x:
         cluster[high(x)]. delete(low(x));
 9:
                                                           delete
                           continued...
```

```
Algorithm 36 delete(x)
                            ...continued
         if cluster[high(x)]. min() = null then
10:
              summary . delete(high(x));
11:
              if x = \max then
12:
13:
                   summax \leftarrow summary.max();
14:
                   if summax = null then max \leftarrow min;
                   else
15:
16:
                        offs \leftarrow cluster[summax]. max();
17:
                        \max \leftarrow summax \circ offs
         else
18:
              if x = \max then
19:
20:
                   offs \leftarrow cluster[high(x)]. max();
                   \max \leftarrow \text{high}(x) \circ \text{offs};
21:
```

```
Algorithm 36 delete(x)
                            ...continued
                                                      fix maximum
         if cluster[high(x)]. min() = null then
10:
              summary . delete(high(x));
11:
              if x = \max then
12:
13:
                   summax \leftarrow summary.max();
                   if summax = null then max \leftarrow min;
14:
                   else
15:
16:
                        offs \leftarrow cluster[summax]. max();
17:
                        \max \leftarrow summax \circ offs
         else
18:
              if x = \max then
19:
20:
                   offs \leftarrow cluster[high(x)]. max();
                   \max \leftarrow \text{high}(x) \circ \text{offs};
21:
```

Note that only one of the possible recusive calls in Line 9 and Line 11 in the deletion-algorithm may take non-constant time.

To see this observe that the call in Line 11 only occurs if the cluster where x was deleted is now empty. But this means that the call in Line 9 deleted the last element in cluster[high(x)]. Such a call only takes constant time.

Hence, we get a recurrence of the form

$$T_{\rm del}(u) = T_{\rm del}(\sqrt{u}) + c$$
.

This gives $T_{del}(u) = \mathcal{O}(\log \log u)$.

7.6 van Emde Boas Trees

Space requirements:

The space requirement fulfills the recurrence

$$S(u) = (\sqrt{u} + 1)S(\sqrt{u}) + \mathcal{O}(\sqrt{u}) .$$

- Note that we cannot solve this recurrence by the Master theorem as the branching factor is not constant.
- One can show by induction that the space requirement is $S(u) = \mathcal{O}(u)$. Exercise.

Let the "real" recurrence relation be

$$S(k^2) = (k+1)S(k) + c_1 \cdot k; S(4) = c_2$$

▶ Replacing S(k) by $R(k) := S(k)/c_2$ gives the recurrence

$$R(k^2) = (k+1)R(k) + ck; R(4) = 1$$

where $c = c_1/c_2 < 1$.

- Now, we show $R(k) \le k 2$ for squares $k \ge 4$.
 - Obviously, this holds for k = 4.
 - For $k = \ell^2 > 4$ with ℓ integral we have

$$R(k) = (1 + \ell)R(\ell) + c\ell$$

$$\leq (1 + \ell)(\ell - 2) + \ell \leq k - 2$$

▶ This shows that R(k) and, hence, S(k) grows linearly.

Dictionary:

- **S.** insert(x): Insert an element x.
- S. delete(x): Delete the element pointed to by x.
- S. search(k): Return a pointer to an element e with key[e] = k in S if it exists; otherwise return null.

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So far we have implemented the search for a key by carefully choosing split-elements.

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Then the memory location of an object x with key k is determined by successively comparing k to split-elements.

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So far we have implemented the search for a key by carefully choosing split-elements.

Then the memory location of an object x with key k is determined by successively comparing k to split-elements.

Hashing tries to directly compute the memory location from the given key. The goal is to have constant search time.

Definitions:

▶ Universe U of keys, e.g., $U \subseteq \mathbb{N}_0$. U very large.

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- ► Hash function $h: U \rightarrow [0, ..., n-1]$.

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The hash-function *h* should fulfill:

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The hash-function h should fulfill:

Fast to evaluate.

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- ► Hash function $h: U \rightarrow [0, ..., n-1]$.

The hash-function h should fulfill:

- Fast to evaluate.
- Small storage requirement.

Definitions:

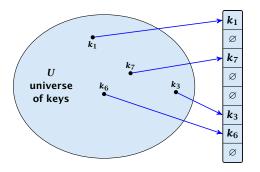
- ▶ Universe U of keys, e.g., $U \subseteq \mathbb{N}_0$. U very large.
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- Array $T[0, \ldots, n-1]$ hash-table.
- ► Hash function $h: U \rightarrow [0, ..., n-1]$.

The hash-function h should fulfill:

- Fast to evaluate.
- Small storage requirement.
- Good distribution of elements over the whole table.

Direct Addressing

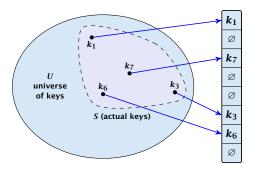
Ideally the hash function maps all keys to different memory locations.



This special case is known as Direct Addressing. It is usually very unrealistic as the universe of keys typically is quite large, and in particular larger than the available memory.

Perfect Hashing

Suppose that we know the set S of actual keys (no insert/no delete). Then we may want to design a simple hash-function that maps all these keys to different memory locations.



Such a hash function h is called a perfect hash function for set S.

If we do not know the keys in advance, the best we can hope for is that the hash function distributes keys evenly across the table.

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7.7 Hashing

Problem: Collisions

Usually the universe U is much larger than the table-size n.

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If we do not know the keys in advance, the best we can hope for is that the hash function distributes keys evenly across the table.

Problem: Collisions

Usually the universe U is much larger than the table-size n.

Hence, there may be two elements k_1, k_2 from the set S that map to the same memory location (i.e., $h(k_1) = h(k_2)$). This is called a collision.



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Lemma 9

The probability of having a collision when hashing m elements into a table of size n under uniform hashing is at least

$$1 - e^{-\frac{m(m-1)}{2n}} \approx 1 - e^{-\frac{m^2}{2n}}$$
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Uniform hashing:

Choose a hash function uniformly at random from all functions $f: U \to [0, ..., n-1]$.



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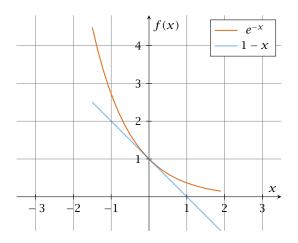
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Here the first equality follows since the ℓ -th element that is hashed has a probability of $\frac{n-\ell+1}{n}$ to not generate a collision under the condition that the previous elements did not induce collisions.

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The inequality $1 - x \le e^{-x}$ is derived by stopping the Taylor-expansion of e^{-x} after the second term.



Resolving Collisions

The methods for dealing with collisions can be classified into the two main types

- open addressing, aka. closed hashing
- hashing with chaining, aka. closed addressing, open hashing.

Resolving Collisions

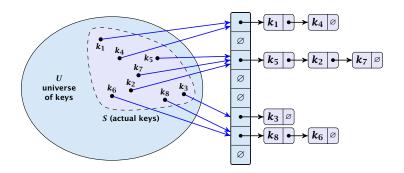
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- open addressing, aka. closed hashing
- hashing with chaining, aka. closed addressing, open hashing.

There are applications e.g. computer chess where you do not resolve collisions at all.

Arrange elements that map to the same position in a linear list.

- Access: compute h(x) and search list for key[x].
- Insert: insert at the front of the list.



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- A⁺ denotes the average time for a successful search when using A;
- ▶ A^- denotes the average time for an **unsuccessful** search when using A;
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$$A^- = 1 + \alpha .$$

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Let for two keys k_i and k_j , X_{ij} denote the indicator variable for the event that k_i and k_j hash to the same position. Clearly, $Pr[X_{i,i} = 1] = 1/n$ for uniform hashing.

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The expected successful search cost is

$$E\left[\frac{1}{m}\sum_{i=1}^{m}\left(1+\sum_{j=i+1}^{m}X_{ij}\right)\right]$$



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$$=1+\frac{1}{mn}\sum_{i=1}^{m}(m-i)$$

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$$= 1+\frac{m-1}{2n}=1+\frac{\alpha}{2}-\frac{\alpha}{2m}.$$

$$\begin{split} \mathbf{E} \left[\frac{1}{m} \sum_{i=1}^{m} \left(1 + \sum_{j=i+1}^{m} X_{ij} \right) \right] &= \frac{1}{m} \sum_{i=1}^{m} \left(1 + \sum_{j=i+1}^{m} \mathbf{E} \left[X_{ij} \right] \right) \\ &= \frac{1}{m} \sum_{i=1}^{m} \left(1 + \sum_{j=i+1}^{m} \frac{1}{n} \right) \\ &= 1 + \frac{1}{mn} \sum_{i=1}^{m} (m-i) \\ &= 1 + \frac{1}{mn} \left(m^2 - \frac{m(m+1)}{2} \right) \\ &= 1 + \frac{m-1}{2n} = 1 + \frac{\alpha}{2} - \frac{\alpha}{2m} \end{split} .$$

Hence, the expected cost for a successful search is $A^+ \leq 1 + \frac{\alpha}{2}$.

Hashing with Chaining

Disadvantages:

- pointers increase memory requirements
- pointers may lead to bad cache efficiency

Advantages:

- no à priori limit on the number of elements
- deletion can be implemented efficiently
- by using balanced trees instead of linked list one can also obtain worst-case guarantees.

All objects are stored in the table itself.

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Define a function h(k, j) that determines the table-position to be examined in the j-th step. The values $h(k, 0), \ldots, h(k, n-1)$ must form a permutation of $0, \ldots, n-1$.

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Search(k): Try position h(k,0); if it is empty your search fails; otw. continue with h(k,1), h(k,2),

Insert(x): Search until you find an empty slot; insert your element there. If your search reaches h(k, n-1), and this slot is non-empty then your table is full.

Choices for h(k, j):

Linear probing:

```
h(k,i) = h(k) + i \mod n
(sometimes: h(k,i) = h(k) + ci \mod n).
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- Quadratic probing: $h(k, i) = h(k) + c_1 i + c_2 i^2 \mod n$.
- Double hashing: $h(k,i) = h_1(k) + ih_2(k) \mod n$.

For quadratic probing and double hashing one has to ensure that the search covers all positions in the table (i.e., for double hashing $h_2(k)$ must be relatively prime to n (teilerfremd); for quadratic probing c_1 and c_2 have to be chosen carefully).

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Linear Probing

Advantage: Cache-efficiency. The new probe position is very likely to be in the cache.

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Lemma 10

Let L be the method of linear probing for resolving collisions:

$$L^+ \approx \frac{1}{2} \left(1 + \frac{1}{1 - \alpha} \right)$$

$$L^- \approx \frac{1}{2} \left(1 + \frac{1}{(1 - \alpha)^2} \right)$$



Quadratic Probing

- Not as cache-efficient as Linear Probing.
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Lemma 11

Let Q be the method of quadratic probing for resolving collisions:

$$Q^+ \approx 1 + \ln\left(\frac{1}{1-\alpha}\right) - \frac{\alpha}{2}$$

$$Q^- \approx \frac{1}{1-\alpha} + \ln\left(\frac{1}{1-\alpha}\right) - \alpha$$

Double Hashing

Any probe into the hash-table usually creates a cache-miss.

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Lemma 12

Let D be the method of double hashing for resolving collisions:

$$D^+ \approx \frac{1}{\alpha} \ln \left(\frac{1}{1 - \alpha} \right)$$

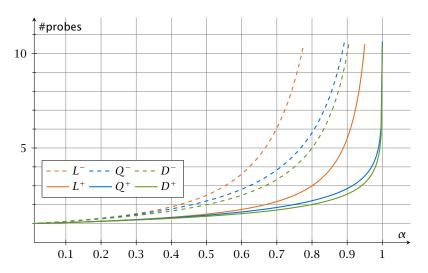
$$D^- \approx \frac{1}{1-\alpha}$$

Some values:

α	Linear Probing		Quadratic Probing		Double Hashing	
	L^+	L^{-}	Q^+	Q^-	D^+	D^-
0.5	1.5	2.5	1.44	2.19	1.39	2
0.9	5.5	50.5	2.85	11.40	2.55	10
0.95	10.5	200.5	3.52	22.05	3.15	20



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We analyze the time for a search in a very idealized Open Addressing scheme.

► The probe sequence h(k,0), h(k,1), h(k,2),... is equally likely to be any permutation of (0,1,...,n-1).

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$$= Pr[A_1] \cdot Pr[A_2 \mid A_1] \cdot Pr[A_3 \mid A_1 \cap A_2] \cdot \dots \cdot Pr[A_{i-1} \mid A_1 \cap \cdots \cap A_{i-2}]$$

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$$\Pr[X \ge i] = \frac{m}{n} \cdot \frac{m-1}{n-1} \cdot \frac{m-2}{n-2} \cdot \dots \cdot \frac{m-i+2}{n-i+2}$$

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$$\le \left(\frac{m}{n}\right)^{i-1}$$

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$$\le \left(\frac{m}{n}\right)^{i-1} = \alpha^{i-1} .$$

E[X]

$$E[X] = \sum_{i=1}^{\infty} \Pr[X \ge i]$$

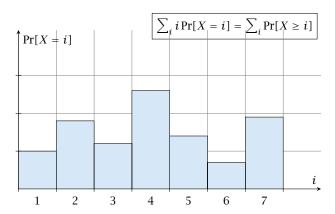
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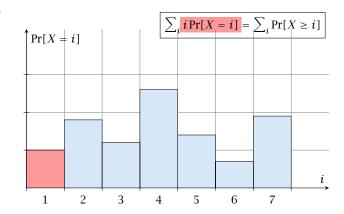
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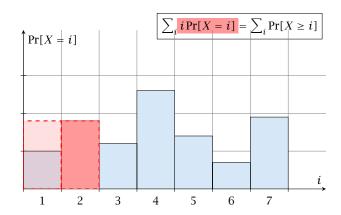
$$\frac{1}{1-\alpha}=1+\alpha+\alpha^2+\alpha^3+\dots$$



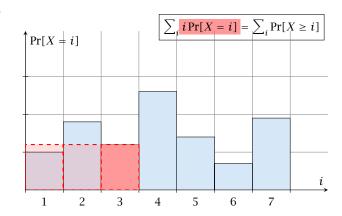
i = 1



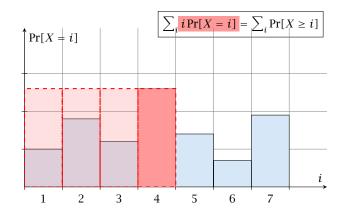
$$i = 2$$



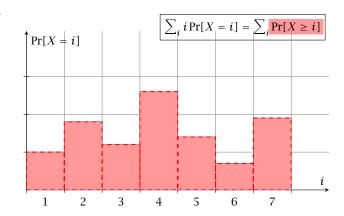
$$i = 3$$



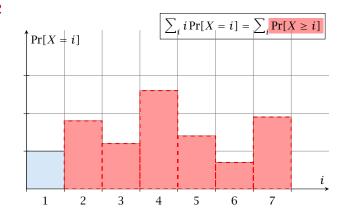
$$i = 4$$



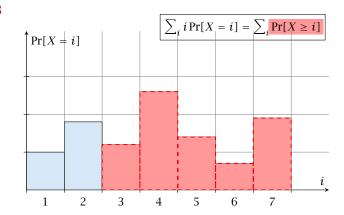
$$i = 1$$



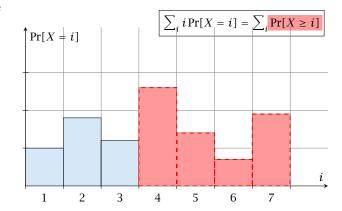
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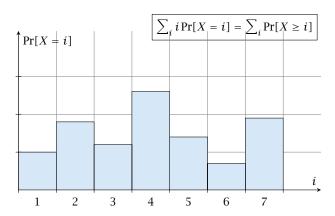


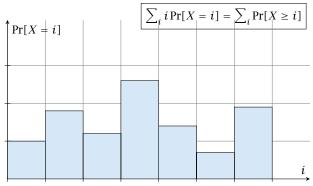
$$i = 3$$



$$i = 4$$







The j-th rectangle² appears in both sums j⁶ times. (j times in the first due to multiplication with j; and j times in the second for summands i = 1, 2, ..., j)

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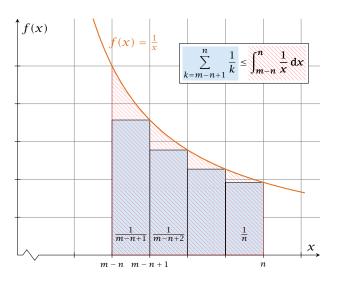
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$$\leq \frac{1}{\alpha} \int_{n-m}^{n} \frac{1}{x} dx = \frac{1}{\alpha} \ln \frac{n}{n-m} = \frac{1}{\alpha} \ln \frac{1}{1-\alpha} .$$

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How do we delete in a hash-table?

For hashing with chaining this is not a problem. Simply search for the key, and delete the item in the corresponding list.

How do we delete in a hash-table?

- For hashing with chaining this is not a problem. Simply search for the key, and delete the item in the corresponding list.
- For open addressing this is difficult.

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- Simply removing a key might interrupt the probe sequence of other keys which then cannot be found anymore.
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 - During an insertion if a deleted-marker is encountered an element can be inserted there.
 - During a search a deleted-marker must not be used to terminate the probe sequence.
- The table could fill up with deleted-markers leading to bad performance.
- ▶ If a table contains many deleted-markers (linear fraction of the keys) one can rehash the whole table and amortize the cost for this rehash against the cost for the deletions.



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For Linear Probing one can delete elements without using deletion-markers.

- For Linear Probing one can delete elements without using deletion-markers.
- Upon a deletion elements that are further down in the probe-sequence may be moved to guarantee that they are still found during a search.

7.7 Hashing

```
Algorithm 37 delete(p)

1: T[p] \leftarrow \text{null}

2: p \leftarrow \text{succ}(p)

3: while T[p] \neq \text{null do}

4: y \leftarrow T[p]

5: T[p] \leftarrow \text{null}

6: p \leftarrow \text{succ}(p)

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p is the index into the table-cell that contains the object to be deleted.

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Pointers into the hash-table become invalid.



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However, the assumption of uniform hashing that h is chosen randomly from all functions $f:U\to [0,\dots,n-1]$ is clearly unrealistic as there are $n^{|U|}$ such functions. Even writing down such a function would take $|U|\log n$ bits.

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However, the assumption of uniform hashing that h is chosen randomly from all functions $f:U\to [0,\dots,n-1]$ is clearly unrealistic as there are $n^{|U|}$ such functions. Even writing down such a function would take $|U|\log n$ bits.

Universal hashing tries to define a set $\mathcal H$ of functions that is much smaller but still leads to good average case behaviour when selecting a hash-function uniformly at random from $\mathcal H$.

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Definition 13

A class $\mathcal H$ of hash-functions from the universe U into the set $\{0,\dots,n-1\}$ is called universal if for all $u_1,u_2\in U$ with $u_1\neq u_2$

$$\Pr[h(u_1) = h(u_2)] \le \frac{1}{n}$$
,

where the probability is w.r.t. the choice of a random hash-function from set \mathcal{H} .

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Note that this means that the probability of a collision between two arbitrary elements is at most $\frac{1}{n}$.

Definition 14

A class \mathcal{H} of hash-functions from the universe U into the set $\{0,\ldots,n-1\}$ is called 2-independent (pairwise independent) if the following two conditions hold

- For any key $u \in U$, and $t \in \{0, ..., n-1\}$ $\Pr[h(u) = t] = \frac{1}{n}$, i.e., a key is distributed uniformly within the hash-table.
- For all $u_1, u_2 \in U$ with $u_1 \neq u_2$, and for any two hash-positions t_1, t_2 :

$$\Pr[h(u_1) = t_1 \land h(u_2) = t_2] \le \frac{1}{n^2} .$$

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.

This requirement clearly implies a universal hash-function.



7.7 Hashing

Definition 15

A class $\mathcal H$ of hash-functions from the universe U into the set $\{0,\ldots,n-1\}$ is called k-independent if for any choice of $\ell \le k$ distinct keys $u_1,\ldots,u_\ell \in U$, and for any set of ℓ not necessarily distinct hash-positions t_1,\ldots,t_ℓ :

$$\Pr[h(u_1) = t_1 \wedge \cdots \wedge h(u_\ell) = t_\ell] \le \frac{1}{n^\ell} ,$$

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Definition 16

A class $\mathcal H$ of hash-functions from the universe U into the set $\{0,\ldots,n-1\}$ is called (μ,k) -independent if for any choice of $\ell \leq k$ distinct keys $u_1,\ldots,u_\ell \in U$, and for any set of ℓ not necessarily distinct hash-positions t_1,\ldots,t_ℓ :

$$\Pr[h(u_1) = t_1 \wedge \cdots \wedge h(u_\ell) = t_\ell] \leq \frac{\mu}{n^\ell} ,$$

where the probability is w.r.t. the choice of a random hash-function from set \mathcal{H} .

7.7 Hashing 6.

Let $U:=\{0,\ldots,p-1\}$ for a prime p. Let $\mathbb{Z}_p:=\{0,\ldots,p-1\}$, and let $\mathbb{Z}_p^*:=\{1,\ldots,p-1\}$ denote the set of invertible elements in \mathbb{Z}_p .

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Define

$$h_{a,b}(x) := (ax + b \bmod p) \bmod n$$

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Define

$$h_{a,b}(x) := (ax + b \bmod p) \bmod n$$

Lemma 17

The class

$$\mathcal{H} = \{ h_{a,b} \mid a \in \mathbb{Z}_p^*, b \in \mathbb{Z}_p \}$$

is a universal class of hash-functions from U to $\{0, \ldots, n-1\}$.

Proof.

Let $x, y \in U$ be two distinct keys. We have to show that the probability of a collision is only 1/n.

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where we use that \mathbb{Z}_p is a field (Körper) and, hence, has no zero divisors (nullteilerfrei).

The hash-function does not generate collisions before the \pmod{n} -operation. Furthermore, every choice (a,b) is mapped to a different pair (t_x,t_y) with $t_x:=ax+b$ and $t_y:=ay+b$.

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$$t_{x} \equiv ax + b \qquad (\text{mod } p)$$

$$t_{y} \equiv ay + b \qquad (\text{mod } p)$$

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$$t_{y} \equiv ay + b \qquad (\text{mod } p)$$

$$t_{x} - t_{y} \equiv a(x - y) \qquad (\text{mod } p)$$

$$t_{y} \equiv ay + b \qquad (\text{mod } p)$$

$$a \equiv (t_{x} - t_{y})(x - y)^{-1} \qquad (\text{mod } p)$$

$$b \equiv t_{y} - ay \qquad (\text{mod } p)$$

There is a one-to-one correspondence between hash-functions (pairs (a, b), $a \ne 0$) and pairs (t_x, t_y) , $t_x \ne t_y$.

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Fix a value t_x . There are p-1 possible values for choosing t_y .

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What happens when we do the mod n operation?

Fix a value t_x . There are p-1 possible values for choosing t_y .

From the range $0, \ldots, p-1$ the values $t_x, t_x + n, t_x + 2n, \ldots$ map to t_x after the modulo-operation. These are at most $\lceil p/n \rceil$ values.

As $t_y \neq t_x$ there are

$$\left\lceil \frac{p}{n} \right\rceil - 1$$

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possibilities for choosing $t_{\mathcal{Y}}$ such that the final hash-value creates a collision.

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possibilities for choosing $t_{\mathcal{Y}}$ such that the final hash-value creates a collision.

This happens with probability at most $\frac{1}{n}$.

It is also possible to show that $\mathcal H$ is an (almost) pairwise independent class of hash-functions.

$$\Pr_{t_{\mathcal{X}} \neq t_{\mathcal{Y}} \in \mathbb{Z}_p^2} \left[\begin{array}{c} t_{\mathcal{X}} \bmod n = h_1 \\ t_{\mathcal{Y}} \bmod n = h_2 \end{array} \right]$$

7.7 Hashing

It is also possible to show that $\mathcal H$ is an (almost) pairwise independent class of hash-functions.

$$\frac{\left\lfloor \frac{p}{n} \right\rfloor^2}{p(p-1)} \le \Pr_{t_X \neq t_Y \in \mathbb{Z}_p^2} \left[\begin{array}{c} t_X \bmod n = h_1 \\ t_Y \bmod n = h_2 \end{array} \right] \le \frac{\left\lceil \frac{p}{n} \right\rceil^2}{p(p-1)}$$

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$$\frac{\left\lfloor \frac{p}{n} \right\rfloor^2}{p(p-1)} \le \Pr_{t_x \neq t_y \in \mathbb{Z}_p^2} \left[\begin{array}{c} t_x \bmod n = h_1 \\ t_y \bmod n = h_2 \end{array} \right] \le \frac{\left\lceil \frac{p}{n} \right\rceil^2}{p(p-1)}$$

Note that the middle is the probability that $h(x) = h_1$ and $h(y) = h_2$. The total number of choices for (t_x, t_y) is p(p-1). The number of choices for t_x (t_y) such that $t_x \bmod n = h_1$ $(t_y \bmod n = h_2)$ lies between $\lfloor \frac{p}{n} \rfloor$ and $\lceil \frac{p}{n} \rceil$.

Definition 18

Let $d \in \mathbb{N}$; $q \ge (d+1)n$ be a prime; and let $\bar{a} \in \{0, \dots, q-1\}^{d+1}$. Define for $x \in \{0, \dots, q-1\}$

$$h_{\bar{a}}(x) := \left(\sum_{i=0}^{d} a_i x^i \bmod q\right) \bmod n$$
.

Let $\mathcal{H}_n^d := \{h_{\bar{a}} \mid \bar{a} \in \{0, \dots, q-1\}^{d+1}\}$. The class \mathcal{H}_n^d is (e, d+1)-independent.

Note that in the previous case we had d = 1 and chose $a_d \neq 0$.

For the coefficients $\bar{a} \in \{0,\ldots,q-1\}^{d+1}$ let $f_{\bar{a}}$ denote the polynomial

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The polynomial is defined by d+1 distinct points.

Fix $\ell \le d+1$; let $x_1, \dots, x_\ell \in \{0, \dots, q-1\}$ be keys, and let t_1, \dots, t_ℓ denote the corresponding hash-function values.

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Then

$$h_{\bar{a}} \in A^{\ell} \Leftrightarrow h_{\bar{a}} = f_{\bar{a}} \bmod n$$
 and

$$f_{\bar{a}}(x_i) \in \underbrace{\{t_i + \alpha \cdot n \mid \alpha \in \{0, \dots, \lceil \frac{q}{n} \rceil - 1\}\}}_{=:B_i}$$

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$$|B_1| \cdot \ldots \cdot |B_{\ell}|$$

possibilities to do this (so that $h_{\bar{a}}(x_i) = t_i$).

Now, we choose $d-\ell+1$ other inputs and choose their value arbitrarily. We have $q^{d-\ell+1}$ possibilities to do this.

Now, we choose $d-\ell+1$ other inputs and choose their value arbitrarily. We have $q^{d-\ell+1}$ possibilities to do this.

Therefore we have

$$|B_1| \cdot \ldots \cdot |B_\ell| \cdot q^{d-\ell+1} \le \lceil \frac{q}{n} \rceil^\ell \cdot q^{d-\ell+1}$$

possibilities to choose \bar{a} such that $h_{\bar{a}} \in A_{\ell}$.

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Therefore the probability of choosing $h_{ ilde{a}}$ from A_{ℓ} is only

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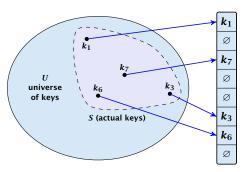
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This shows that the \mathcal{H} is (e, d+1)-universal.

The last step followed from $q \ge (d+1)n$, and $\ell \le d+1$.

Suppose that we **know** the set S of actual keys (no insert/no delete). Then we may want to design a **simple** hash-function that maps all these keys to different memory locations.



7.7 Hashing

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Can we get an upper bound on the probability of having collisions?

The probability of having 1 or more collisions can be at most $\frac{1}{2}$ as otherwise the expectation would be larger than $\frac{1}{2}$.



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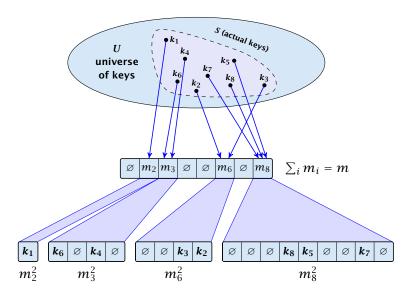
We construct a two-level scheme. We first use a hash-function that maps elements from ${\cal S}$ to ${\cal m}$ buckets.

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However, a hash-table size of $n = m^2$ is very very high.

We construct a two-level scheme. We first use a hash-function that maps elements from $\mathcal S$ to m buckets.

Let m_j denote the number of items that are hashed to the j-th bucket. For each bucket we choose a second hash-function that maps the elements of the bucket into a table of size m_j^2 . The second function can be chosen such that all elements are mapped to different locations.





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The first expectation is simply the expected number of collisions, for the first level. Since we use universal hashing we have

$$= 2\binom{m}{2} \frac{1}{m} + m = 2m - 1 .$$

We need only $\mathcal{O}(m)$ time to construct a hash-function h with $\sum_j m_j^2 = \mathcal{O}(4m)$, because with probability at least 1/2 a random function from a universal family will have this property.

Then we construct a hash-table h_j for every bucket. This takes expected time $\mathcal{O}(m_j)$ for every bucket. A random function h_j is collision-free with probability at least 1/2. We need $\mathcal{O}(m_j)$ to test this.

We only need that the hash-functions are chosen from a universal family!!!

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Goal:

Try to generate a hash-table with constant worst-case search time in a dynamic scenario.

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Try to generate a hash-table with constant worst-case search time in a dynamic scenario.

- ▶ Two hash-tables $T_1[0,...,n-1]$ and $T_2[0,...,n-1]$, with hash-functions h_1 , and h_2 .
- ▶ An object x is either stored at location $T_1[h_1(x)]$ or $T_2[h_2(x)]$.

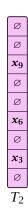
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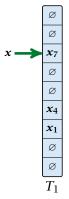
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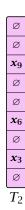
- ▶ Two hash-tables $T_1[0,...,n-1]$ and $T_2[0,...,n-1]$, with hash-functions h_1 , and h_2 .
- An object x is either stored at location $T_1[h_1(x)]$ or $T_2[h_2(x)]$.
- A search clearly takes constant time if the above constraint is met.

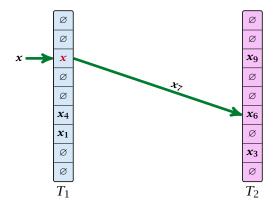
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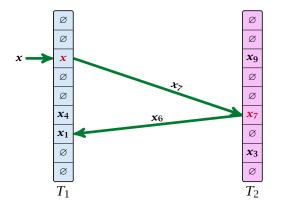


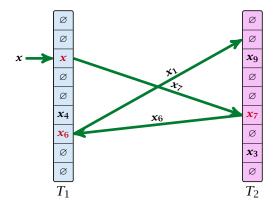












Algorithm 38 Cuckoo-Insert(x)

```
1: if T_1[h_1(x)] = x \vee T_2[h_2(x)] = x then return

2: steps \leftarrow 1

3: while steps \leq maxsteps do

4: exchange x and T_1[h_1(x)]

5: if x = \text{null} then return

6: exchange x and T_2[h_2(x)]

7: if x = \text{null} then return

8: steps \leftarrow steps +1

9: rehash() // change hash-functions; rehash everything

10: Cuckoo-Insert(x)
```

► We call one iteration through the while-loop a step of the algorithm.

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- We call a sequence of iterations through the while-loop without the termination condition becoming true a phase of the algorithm.
- We say a phase is successful if it is not terminated by the maxstep-condition, but the while loop is left because x = null.

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What is the expected time for an insert-operation?

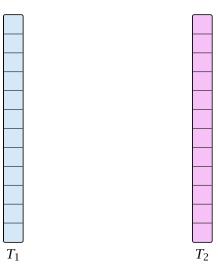
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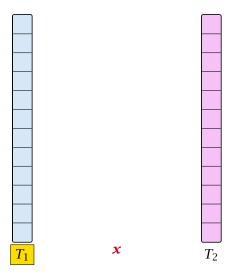
We first analyze the probability that we end-up in an infinite loop (that is then terminated after maxsteps steps).

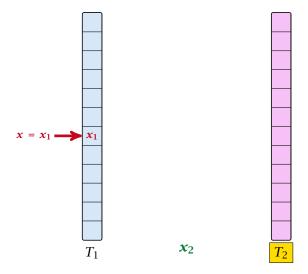
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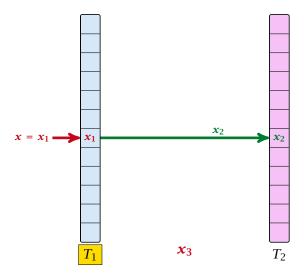
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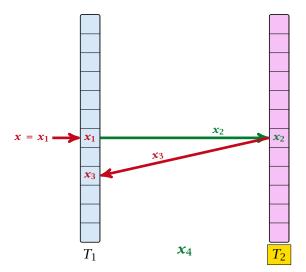
Formally what is the probability to enter an infinite loop that touches s different keys?

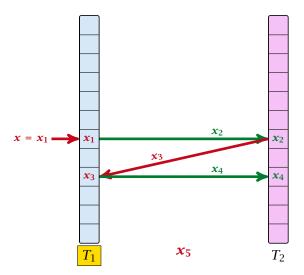


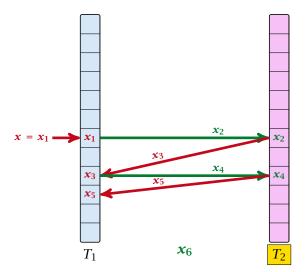


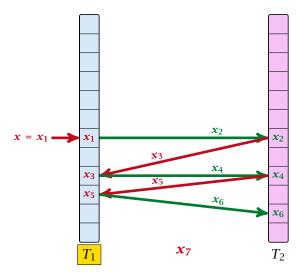




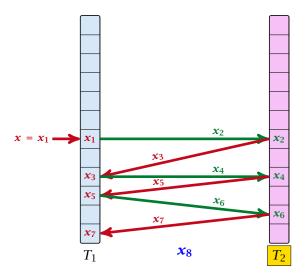


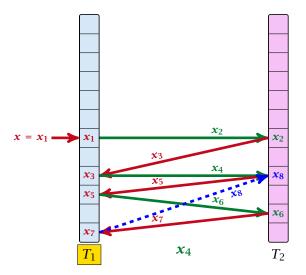


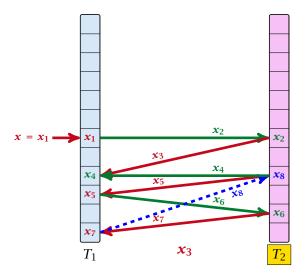


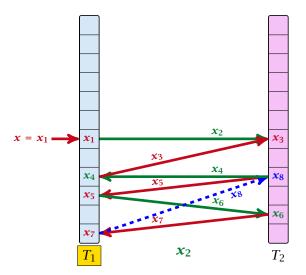


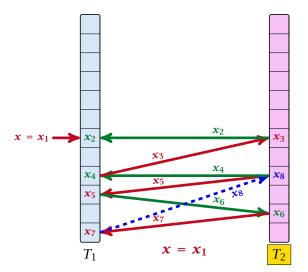


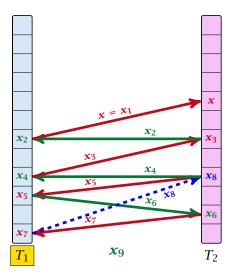


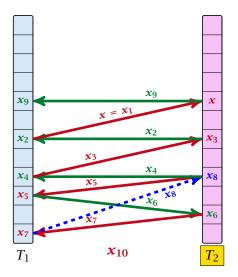


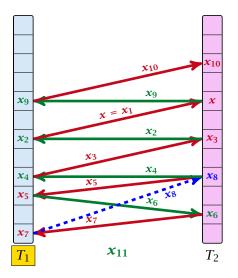


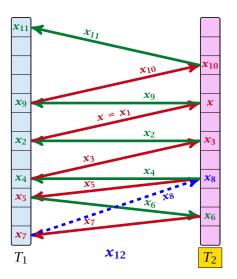


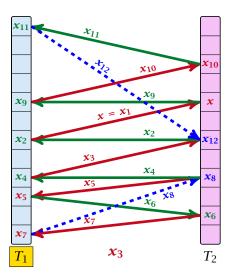


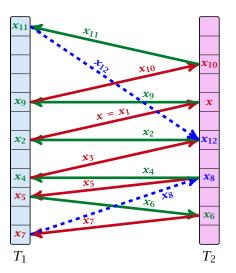


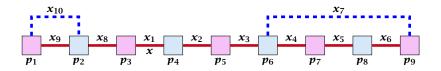






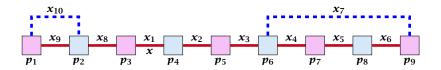






A cycle-structure of size s is defined by

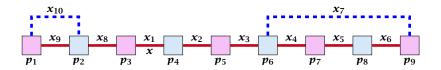
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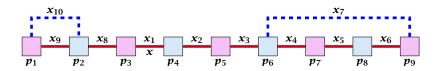
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A cycle-structure of size s is defined by

- ▶ s-1 different cells (alternating btw. cells from T_1 and T_2).
- ▶ *s* distinct keys $x = x_1, x_2, ..., x_s$, linking the cells.

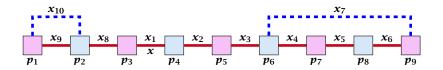
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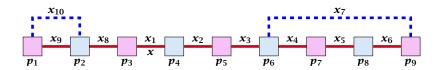
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- The leftmost cell is "linked forward" to some cell on the right.
- The rightmost cell is "linked backward" to a cell on the left.
- ▶ One link represents key x; this is where the counting starts.



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A cycle-structure is active if for every key x_{ℓ} (linking a cell p_i from T_1 and a cell p_j from T_2) we have

$$h_1(x_\ell) = p_i$$
 and $h_2(x_\ell) = p_j$

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Observation:

If during a phase the insert-procedure runs into a cycle there must exist an active cycle structure of size $s \ge 3$.

What is the probability that all keys in a cycle-structure of size s correctly map into their T_1 -cell?

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This probability is at most $\frac{\mu}{n^s}$ since h_1 is a (μ,s) -independent hash-function.

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These events are independent.

The probability that a given cycle-structure of size s is active is at most $\frac{\mu^2}{n^{2s}}$.

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What is the probability that there exists an active cycle structure of size *s*?

The number of cycle-structures of size s is at most

$$s^3 \cdot n^{s-1} \cdot m^{s-1}$$
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- ▶ There are n^{s-1} possibilities to choose the cells.

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The probability that there exists an active cycle-structure is therefore at most

$$\sum_{s=3}^{\infty} s^3 \cdot n^{s-1} \cdot m^{s-1} \cdot \frac{\mu^2}{n^{2s}}$$

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$$\leq \frac{\mu^2}{m^2} \sum_{s=3}^{\infty} s^3 \left(\frac{1}{1+\epsilon}\right)^s$$

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The probability that there exists an active cycle-structure is therefore at most

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Here we used the fact that $(1 + \epsilon)m \le n$.

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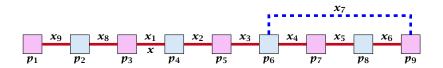
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Here we used the fact that $(1 + \epsilon)m \le n$.

Hence,

$$\Pr[\mathsf{cycle}] = \mathcal{O}\left(\frac{1}{m^2}\right)$$
.

Now, we analyze the probability that a phase is not successful without running into a closed cycle.



Sequence of visited keys:

$$x = x_1, x_2, x_3, x_4, x_5, x_6, x_7, x_3, x_2, x_1 = x, x_8, x_9, \dots$$



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Consider the sequence of not necessarily distinct keys starting with x in the order that they are visited during the phase.

Consider the sequence of not necessarily distinct keys starting with \boldsymbol{x} in the order that they are visited during the phase.

Lemma 19

If the sequence is of length p then there exists a sub-sequence of at least $\frac{p+2}{3}$ keys starting with x of distinct keys.

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Proof.

Let i be the number of keys (including x) that we see before the first repeated key. Let j denote the total number of distinct keys.

The sequence is of the form:

$$x = x_1 \rightarrow x_2 \rightarrow \cdots \rightarrow x_i \rightarrow x_r \rightarrow x_{r-1} \rightarrow \cdots \rightarrow x_1 \rightarrow x_{i+1} \rightarrow \cdots \rightarrow x_j$$

As $r \le i - 1$ the length p of the sequence is

$$p = i + r + (j - i) \le i + j - 1$$
.

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Proof.

Let i be the number of keys (including x) that we see before the first repeated key. Let j denote the total number of distinct keys.

The sequence is of the form:

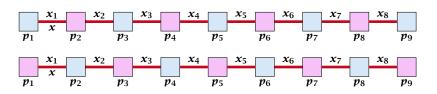
$$x = x_1 \rightarrow x_2 \rightarrow \cdots \rightarrow x_i \rightarrow x_r \rightarrow x_{r-1} \rightarrow \cdots \rightarrow x_1 \rightarrow x_{i+1} \rightarrow \cdots \rightarrow x_j$$

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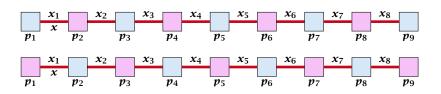
Either sub-sequence $x_1 \rightarrow x_2 \rightarrow \cdots \rightarrow x_i$ or sub-sequence $x_1 \rightarrow x_{i+1} \rightarrow \cdots \rightarrow x_j$ has at least $\frac{p+2}{3}$ elements.





A path-structure of size s is defined by

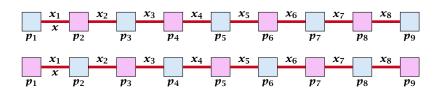
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A path-structure of size s is defined by

ightharpoonup s+1 different cells (alternating btw. cells from T_1 and T_2).

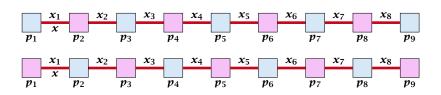
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- ightharpoonup s + 1 different cells (alternating btw. cells from T_1 and T_2).
- s distinct keys $x = x_1, x_2, ..., x_s$, linking the cells.
- ▶ The leftmost cell is either from T_1 or T_2 .

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A path-structure is active if for every key x_ℓ (linking a cell p_i from T_1 and a cell p_j from T_2) we have

$$h_1(x_\ell) = p_i$$
 and $h_2(x_\ell) = p_j$

Observation:

If a phase takes at least t steps without running into a cycle there must exist an active path-structure of size (2t+2)/3.

The probability that a given path-structure of size s is active is at most $\frac{\mu^2}{n^{2s}}$.

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The probability that there exists an active path-structure of size \boldsymbol{s} is at most

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$$\leq 2\mu^2 \left(\frac{m}{n}\right)^{s-1}$$

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Plugging in s = (2t + 2)/3 gives

$$\leq 2\mu^2 \left(\frac{1}{1+\epsilon}\right)^{(2t+2)/3-1} = 2\mu^2 \left(\frac{1}{1+\epsilon}\right)^{(2t-1)/3} \ .$$

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Pr[unsuccessful | no cycle]

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Pr[unsuccessful | no cycle] \leq \Pr[\exists \text{ active path-structure of size at least } \frac{2\text{maxsteps}+2}{3}] \leq \Pr[\exists \text{ active path-structure of size at least } \ell+1] \leq \Pr[\exists \text{ active path-structure of size exactly } \ell+1]
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\begin{split} &\Pr[\mathsf{unsuccessful} \mid \mathsf{no} \; \mathsf{cycle}] \\ &\leq \Pr[\exists \; \mathsf{active} \; \mathsf{path\text{-}structure} \; \mathsf{of} \; \mathsf{size} \; \mathsf{at} \; \mathsf{least} \; \frac{2\mathsf{maxsteps} + 2}{3}] \\ &\leq \Pr[\exists \; \mathsf{active} \; \mathsf{path\text{-}structure} \; \mathsf{of} \; \mathsf{size} \; \mathsf{at} \; \mathsf{least} \; \ell + 1] \\ &\leq \Pr[\exists \; \mathsf{active} \; \mathsf{path\text{-}structure} \; \mathsf{of} \; \mathsf{size} \; \mathsf{exactly} \; \ell + 1] \\ &\leq 2\mu^2 \Big(\frac{1}{1+\epsilon}\Big)^\ell \end{split}
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This gives maxsteps = $\Theta(\log m)$.

So far we estimated

$$\Pr[\mathsf{cycle}] \leq \mathcal{O}\left(\frac{1}{m^2}\right)$$

and

$$\Pr[\mathsf{unsuccessful} \mid \mathsf{no} \; \mathsf{cycle}] \leq \mathcal{O}(\frac{1}{m^2})$$

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Pr[successful] = Pr[no cycle] - Pr[unsuccessful | no cycle]

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We have

Pr[search at least t steps | successful]

= $Pr[search at least t steps \land successful] / Pr[successful]$

The expected number of complete steps in the successful phase of an insert operation is:

$$\begin{aligned} & \text{E}[\text{number of steps} \mid \text{phase successful}] \\ &= \sum_{t \geq 1} \Pr[\text{search takes at least } t \text{ steps} \mid \text{phase successful}] \end{aligned}$$

We have

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\begin{split} \Pr[\mathsf{search} \ \mathsf{at} \ \mathsf{least} \ t \ \mathsf{steps} \ | \ \mathsf{successful}] \\ &= \Pr[\mathsf{search} \ \mathsf{at} \ \mathsf{least} \ t \ \mathsf{steps} \ \land \ \mathsf{successful}] / \Pr[\mathsf{successful}] \\ &\leq \frac{1}{c} \Pr[\mathsf{search} \ \mathsf{at} \ \mathsf{least} \ t \ \mathsf{steps} \ \land \ \mathsf{successful}] / \Pr[\mathsf{no} \ \mathsf{cycle}] \end{split}
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Hence,

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Hence,

E[number of steps | phase successful]

$$\leq \frac{1}{c} \sum_{t \geq 1} \Pr[\text{search at least } t \text{ steps} \mid \text{no cycle}]$$

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This means the expected cost for a successful phase is constant (even after accounting for the cost of the incomplete step that finishes the phase).

A phase that is not successful induces cost for doing a complete rehash (this dominates the cost for the steps in the phase).

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The probability that a phase is not successful is $q = O(1/m^2)$ (probability $\mathcal{O}(1/m^2)$ of running into a cycle and probability $\mathcal{O}(1/m^2)$ of reaching maxsteps without running into a cycle).

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A rehash try requires m insertions and takes expected constant time per insertion. It fails with probability p := O(1/m).

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The expected number of unsuccessful rehashes is $\sum_{i\geq 1} p^i = \frac{1}{1-p} - 1 = \frac{p}{1-p} = \mathcal{O}(p).$



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A phase that is not successful induces cost for doing a complete rehash (this dominates the cost for the steps in the phase).

The probability that a phase is not successful is $q = \mathcal{O}(1/m^2)$ (probability $\mathcal{O}(1/m^2)$ of running into a cycle and probability $\mathcal{O}(1/m^2)$ of reaching maxsteps without running into a cycle).

A rehash try requires m insertions and takes expected constant time per insertion. It fails with probability p := O(1/m).

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Therefore the expected cost for re-hashes is $\mathcal{O}(m) \cdot \mathcal{O}(p) = \mathcal{O}(1)$.



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Let Y_i denote the event that the i-th rehash occurs and does not lead to a valid configuration (i.e., one of the m+1 insertions fails):

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$$\Pr[Y_i|Z_i] \le (m+1) \cdot \mathcal{O}(1/m^2) \le \mathcal{O}(1/m) =: p.$$

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$$\Pr[Z_i] \le \Pr[\wedge_{j=0}^{i-1} Y_j] \le p^i$$

Let Y_i denote the event that the *i*-th rehash occurs and does not lead to a valid configuration (i.e., one of the m+1 insertions fails):

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Let Z_i denote the event that the *i*-th rehash occurs:

The 0-th (re)hash is the initial configuration when doing the $\Pr[Z_i] \leq \Pr[\wedge_{j=0}^{i-1} Y_j] \leq p^i$ insert.

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Let X_i^s , $s \in \{1, ..., m+1\}$ denote the cost for inserting the s-th element during the *i*-th rehash (assuming *i*-th rehash occurs):

$$E[X_i^s]$$

Let Y_i denote the event that the *i*-th rehash occurs and does not lead to a valid configuration (i.e., one of the m+1 insertions fails):

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$$\begin{split} \mathbf{E}[X_i^s] &= \mathbf{E}[\mathsf{steps} \mid \mathsf{phase} \; \mathsf{successful}] \cdot \Pr[\mathsf{phase} \; \mathsf{sucessful}] \\ &+ \mathsf{maxsteps} \cdot \Pr[\mathsf{not} \; \mathsf{sucessful}] \end{split}$$

Let Y_i denote the event that the *i*-th rehash occurs and does not lead to a valid configuration (i.e., one of the m+1 insertions fails):

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$$\mathrm{E}\left[\sum_{i}\sum_{s}Z_{i}X_{i}^{s}\right]$$

$$E\left[\sum_{i}\sum_{s}Z_{i}X_{i}^{s}\right]$$

$$E\left[\sum_{i}\sum_{s}Z_{i}X_{s}^{i}\right] = \sum_{i}\sum_{s}E[Z_{i}] \cdot E[X_{s}^{i}]$$

$$E\left[\sum_{i}\sum_{s}Z_{i}X_{i}^{s}\right]$$

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$$\leq \mathcal{O}(m) \cdot \sum_{i}p^{i}$$

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Therefore, it is sufficient to have $(\mu, \Theta(\log m))$ -independent hash-functions.

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- Therefore we can amortize the rehash cost after a change in table-size against the cost for insertions and deletions.



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Lemma 20

Cuckoo Hashing has an expected constant insert-time and a worst-case constant search-time.

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Note that the above lemma only holds if the fill-factor (number of keys/total number of hash-table slots) is at most $\frac{1}{2(1+\epsilon)}$.

The $1/(2(1+\epsilon))$ fill-factor comes from the fact that the total hash-table is of size 2n (because we have two tables of size n); moreover $m \le (1+\epsilon)n$.

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Sometimes we also have

 \triangleright S. merge(S'): $S := S \cup S'$; $S' := \emptyset$.



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- S. decrease-key(h, k): Decreases the key of the element specified by handle h to k. Assumes that the key is at least k before the operation.

Dijkstra's Shortest Path Algorithm

```
Algorithm 1 Shortest-Path(G = (V, E, d), s \in V)
 1: Input: weighted graph G = (V, E, d); start vertex s;
 2: Output: key-field of every node contains distance from s;
 3: S.build(); // build empty priority queue
 4: for all v \in V \setminus \{s\} do
 5: v \cdot \text{key} \leftarrow \infty;
 6: h_v \leftarrow S.insert(v);
 7: s. \text{key} \leftarrow 0; S. \text{insert}(s);
 8: while S.is-empty() = false do
 9:
     v \leftarrow S. \mathsf{delete\text{-}min}():
10: for all x \in V s.t. (v, x) \in E do
11:
                 if x. key > v. key +d(v,x) then
12:
                       S.decrease-key(h_x, v. key + d(v, x));
13:
                       x. \text{key} \leftarrow v. \text{key} + d(v, x);
```

Prim's Minimum Spanning Tree Algorithm

```
Algorithm 2 Prim-MST(G = (V, E, d), s \in V)
 1: Input: weighted graph G = (V, E, d); start vertex s;
 2: Output: pred-fields encode MST;
 3: S.build(); // build empty priority queue
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14:
                      x. pred \leftarrow v:
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Analysis of Dijkstra and Prim

Both algorithms require:

- 1 build() operation
- ightharpoonup |V| insert() operations
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How good a running time can we obtain?

Operation	Binary Heap	BST	Binomial Heap	Fibonacci Heap*
build	n	$n \log n$	$n \log n$	n
minimum	1	$\log n$	$\log n$	1
is-empty	1	1	1	1
insert	$\log n$	$\log n$	$\log n$	1
delete	$\log n^{**}$	$\log n$	$\log n$	$\log n$
delete-min	$\log n$	$\log n$	$\log n$	$\log n$
decrease-key	$\log n$	$\log n$	$\log n$	1
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Note that most applications use $\mathbf{build}()$ only to create an empty heap which then costs time 1.

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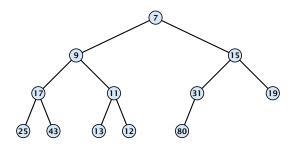
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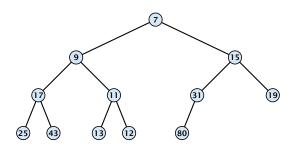
Fibonacci heaps only give an amortized guarantee.

Using Binary Heaps, Prim and Dijkstra run in time $\mathcal{O}((|V|+|E|)\log |V|)$.

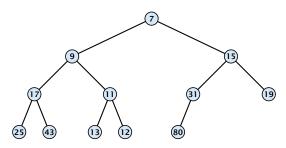
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Nearly complete binary tree; only the last level is not full, and this one is filled from left to right.



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- Heap property: A node's key is not larger than the key of one of its children.



Binary Heaps

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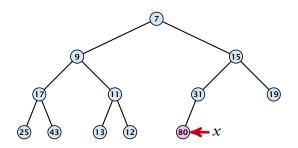
minimum(): return the root-element. Time O(1).

Binary Heaps

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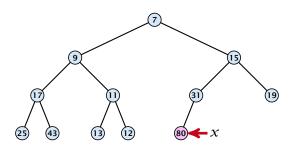
- **minimum()**: return the root-element. Time $\mathcal{O}(1)$.
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Maintain a pointer to the last element x.



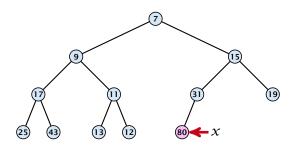
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We can compute the predecessor of x (last element when x is deleted) in time $\mathcal{O}(\log n)$.



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We can compute the predecessor of x (last element when x is deleted) in time O(log n). go up until the last edge used was a right edge. go left; go right until you reach a leaf

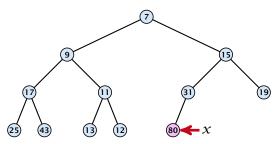


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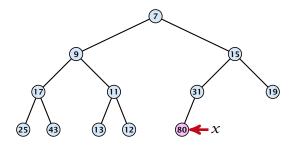
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if you hit the root on the way up, go to the rightmost element

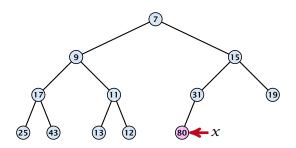


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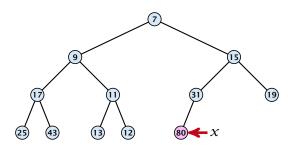
Maintain a pointer to the last element x.

• We can compute the successor of x (last element when an element is inserted) in time $\mathcal{O}(\log n)$.



Maintain a pointer to the last element x.

We can compute the successor of *x* (last element when an element is inserted) in time *O* (log *n*).
 go up until the last edge used was a left edge.
 go right; go left until you reach a null-pointer.

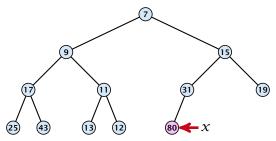


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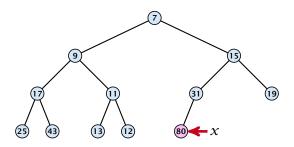
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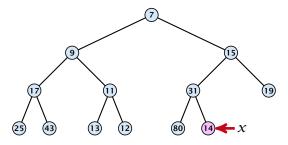
if you hit the root on the way up, go to the leftmost element; insert a new element as a left child;



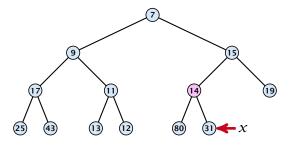
1. Insert element at successor of x.



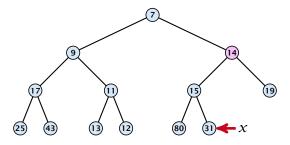
- 1. Insert element at successor of x.
- 2. Exchange with parent until heap property is fulfilled.



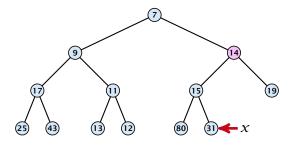
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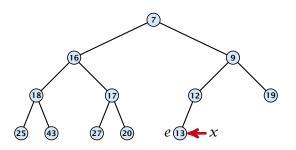


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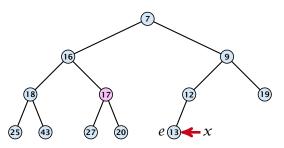


Note that an exchange can either be done by moving the data or by changing pointers. The latter method leads to an addressable priority queue.

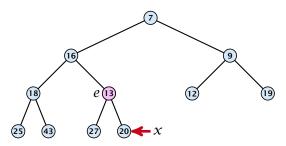
1. Exchange the element to be deleted with the element e pointed to by x.



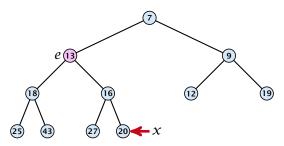
- 1. Exchange the element to be deleted with the element e pointed to by x.
- **2.** Restore the heap-property for the element e.



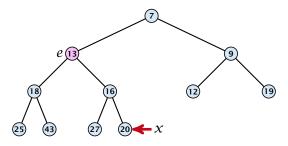
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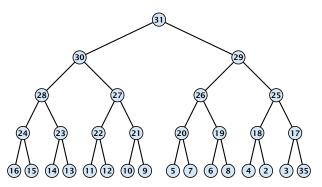


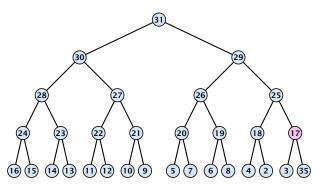
At its new position e may either travel up or down in the tree (but not both directions).

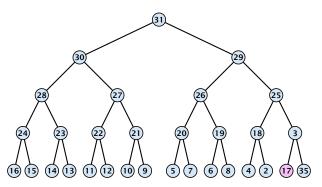
Binary Heaps

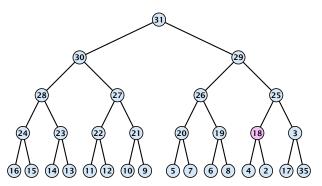
Operations:

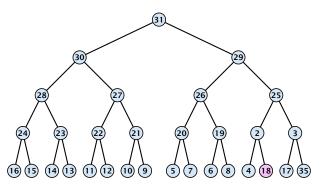
- **minimum()**: return the root-element. Time O(1).
- **is-empty():** check whether root-pointer is null. Time O(1).
- insert(k): insert at successor of x and bubble up. Time $O(\log n)$.
- **delete**(h): swap with x and bubble up or sift-down. Time $O(\log n)$.

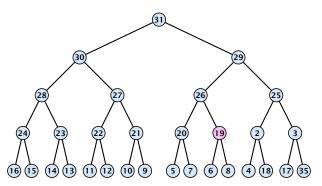


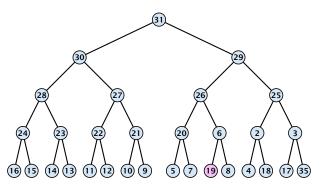


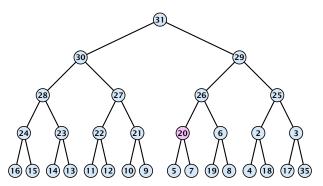


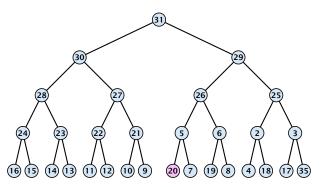


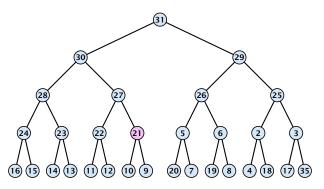


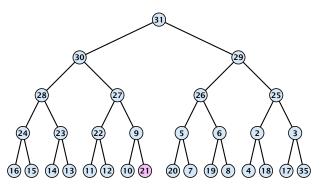


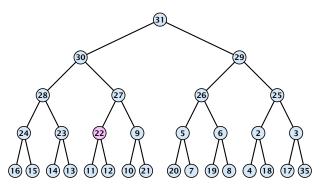


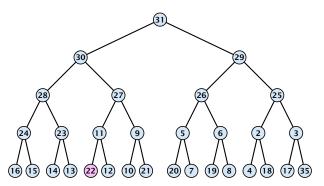


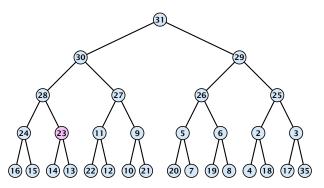


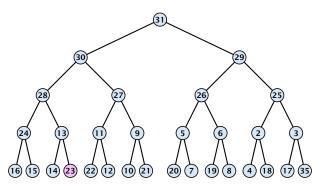


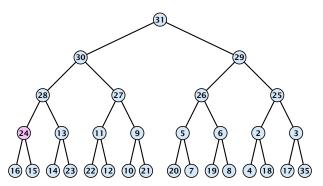


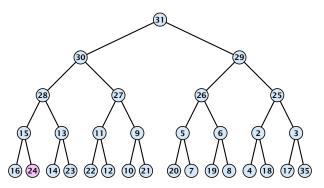


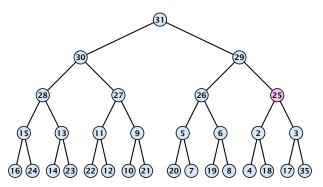


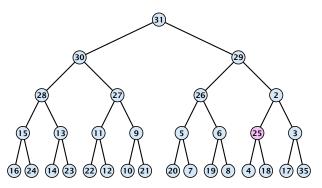


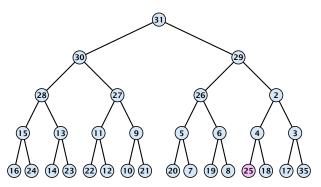


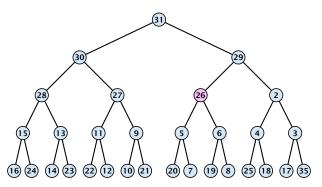


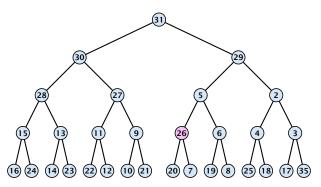


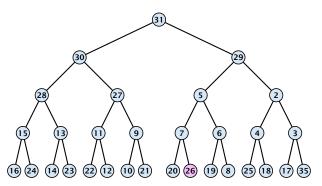


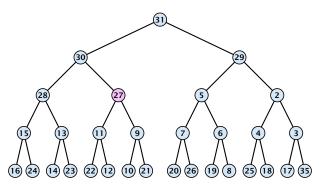


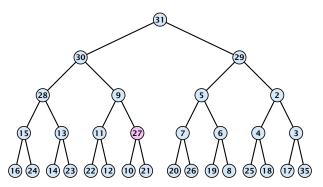


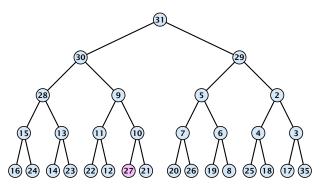


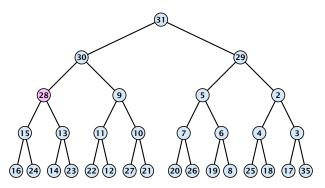


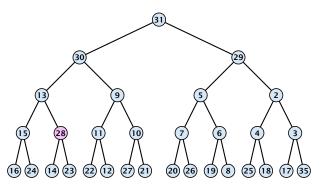


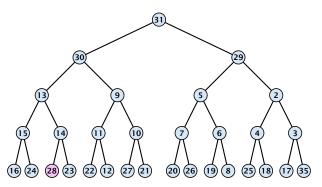


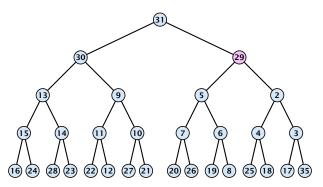


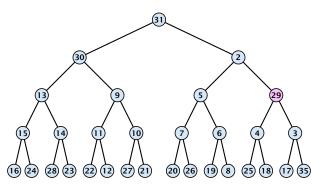


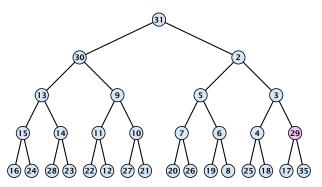


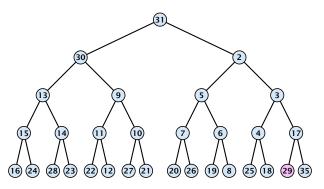


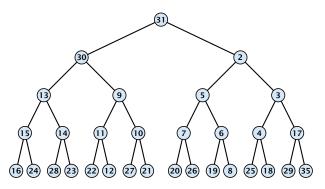


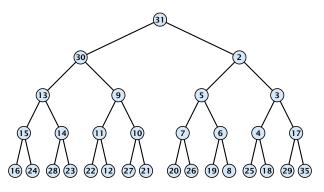












$$\sum_{\text{levels }\ell} 2^{\ell} \cdot (h - \ell) = \sum_{i} i 2^{h-i} = \mathcal{O}(2^h) = \mathcal{O}(n)$$

Operations:

- **minimum():** Return the root-element. Time $\mathcal{O}(1)$.
- **is-empty():** Check whether root-pointer is null. Time $\mathcal{O}(1)$.
- ▶ insert(k): Insert at x and bubble up. Time O(log n).
- **delete**(h): Swap with x and bubble up or sift-down. Time $O(\log n)$.
- **build** (x_1, \ldots, x_n) : Insert elements arbitrarily; then do sift-down operations starting with the lowest layer in the tree. Time $\mathcal{O}(n)$.

The standard implementation of binary heaps is via arrays. Let A[0,...,n-1] be an array

- ▶ The parent of i-th element is at position $\lfloor \frac{i-1}{2} \rfloor$.
- ▶ The left child of i-th element is at position 2i + 1.
- ► The right child of i-th element is at position 2i + 2.

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Finding the successor of x is much easier than in the description on the previous slide. Simply increase or decrease x.

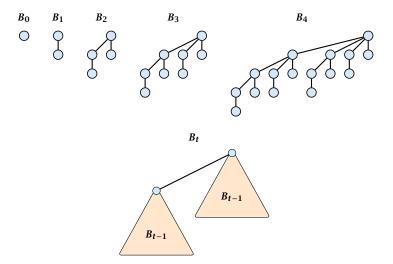
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The resulting binary heap is not addressable. The elements don't maintain their positions and therefore there are no stable handles.

Operation	Binary Heap	BST	Binomial Heap	Fibonacci Heap*
build	n	$n \log n$	$n \log n$	n
minimum	1	$\log n$	$\log n$	1
is-empty	1	1	1	1
insert	$\log n$	$\log n$	$\log n$	1
delete	$\log n^{**}$	$\log n$	$\log n$	$\log n$
delete-min	$\log n$	$\log n$	$\log n$	$\log n$
decrease-key	$\log n$	$\log n$	$\log n$	1
merge	n	$n \log n$	$\log n$	1



Properties of Binomial Trees

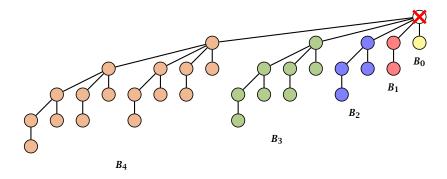
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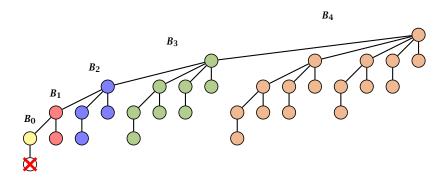
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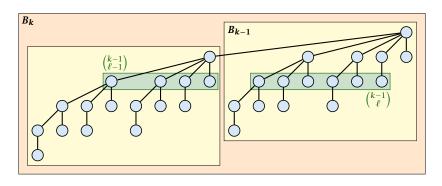
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- $ightharpoonup B_k$ has height k.
- ▶ The root of B_k has degree k.
- ▶ B_k has $\binom{k}{\ell}$ nodes on level ℓ .
- ▶ Deleting the root of B_k gives trees B_0, B_1, \dots, B_{k-1} .



Deleting the root of B_5 leaves sub-trees B_4 , B_3 , B_2 , B_1 , and B_0 .



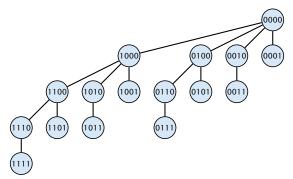
Deleting the leaf furthest from the root (in B_5) leaves a path that connects the roots of sub-trees B_4 , B_3 , B_2 , B_1 , and B_0 .

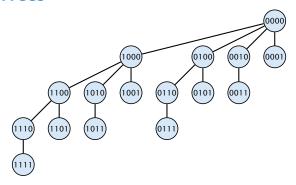


The number of nodes on level ℓ in tree B_k is therefore

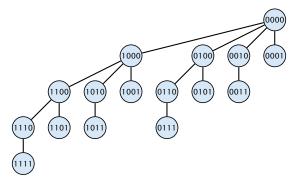
$$\begin{pmatrix} k-1\\\ell-1 \end{pmatrix} + \begin{pmatrix} k-1\\\ell \end{pmatrix} = \begin{pmatrix} k\\\ell \end{pmatrix}$$





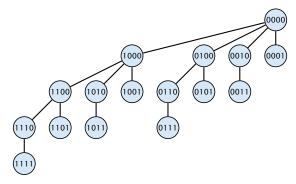


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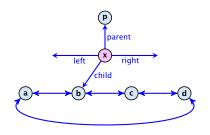
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The ℓ -th level contains nodes that have ℓ 1's in their label.



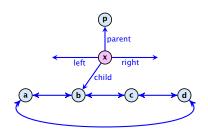
How do we implement trees with non-constant degree?

The children of a node are arranged in a circular linked list.



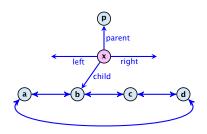
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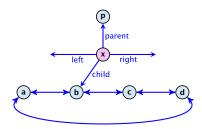
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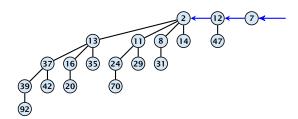


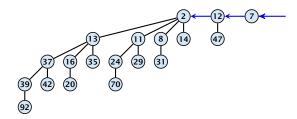
How do we implement trees with non-constant degree?

- The children of a node are arranged in a circular linked list.
- A child-pointer points to an arbitrary node within the list.
- A parent-pointer points to the parent node.
- Pointers x. left and x. right point to the left and right sibling of x (if x does not have siblings then x. left = x. right = x).

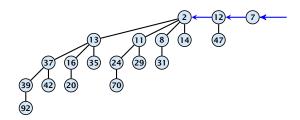


- \blacktriangleright Given a pointer to a node x we can splice out the sub-tree rooted at x in constant time.
- We can add a child-tree T to a node x in constant time if we are given a pointer to x and a pointer to the root of T.



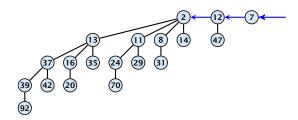


In a binomial heap the keys are arranged in a collection of binomial trees.



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Every tree fulfills the heap-property



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Every tree fulfills the heap-property

There is at most one tree for every dimension/order. For example the above heap contains trees B_0 , B_1 , and B_4 .

Binomial Heap: Merge

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Given the number n of keys to be stored in a binomial heap we can deduce the binomial trees that will be contained in the collection.

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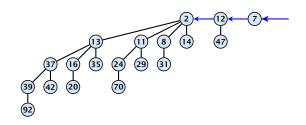
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Let B_{k_1} , B_{k_2} , B_{k_3} , $k_i < k_{i+1}$ denote the binomial trees in the collection and recall that every tree may be contained at most once.

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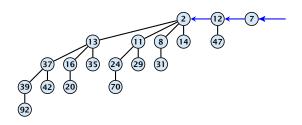
Let B_{k_1} , B_{k_2} , B_{k_3} , $k_i < k_{i+1}$ denote the binomial trees in the collection and recall that every tree may be contained at most once.

Then $n=\sum_i 2^{k_i}$ must hold. But since the k_i are all distinct this means that the k_i define the non-zero bit-positions in the binary representation of n.

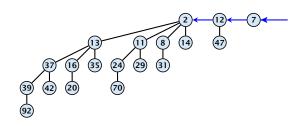


Properties of a heap with n keys:

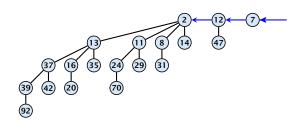
Let $n = b_d b_{d-1}, \dots, b_0$ denote binary representation of n.



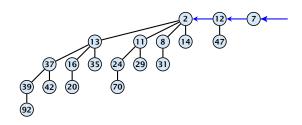
- Let $n = b_d b_{d-1}, \dots, b_0$ denote binary representation of n.
- ▶ The heap contains tree B_i iff $b_i = 1$.



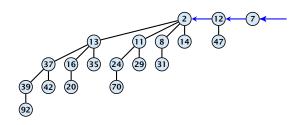
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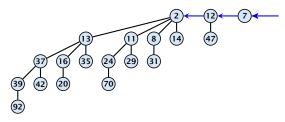
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- ► Hence, at most $\lfloor \log n \rfloor + 1$ trees.
- The minimum must be contained in one of the roots.
- ▶ The height of the largest tree is at most $\lfloor \log n \rfloor$.
- The trees are stored in a single-linked list; ordered by dimension/size.



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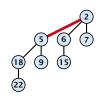
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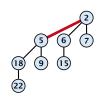
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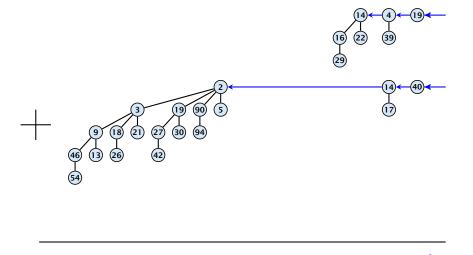
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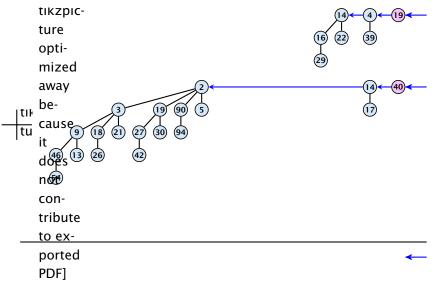
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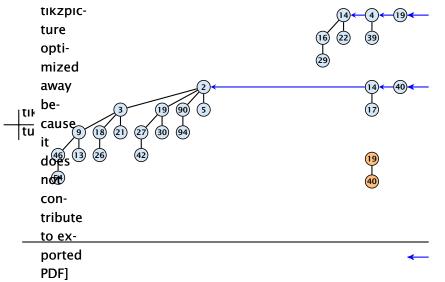
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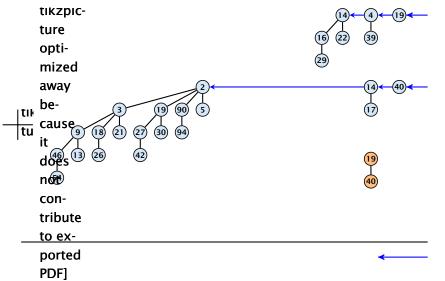
For more trees the technique is analogous to binary addition.

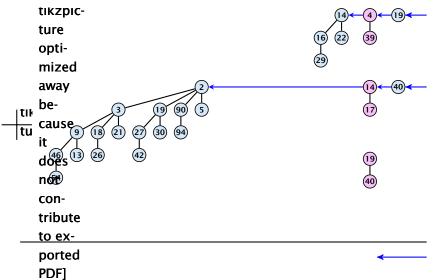


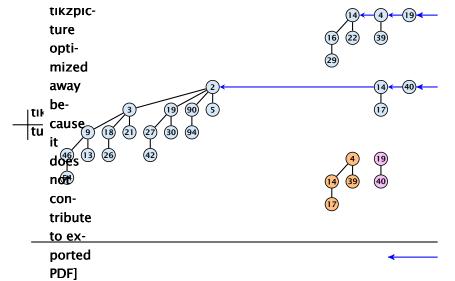


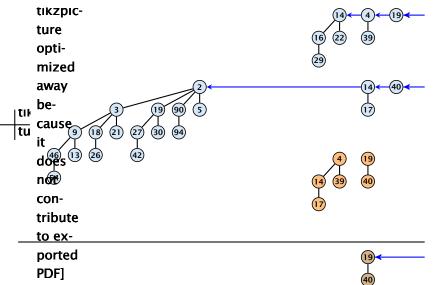


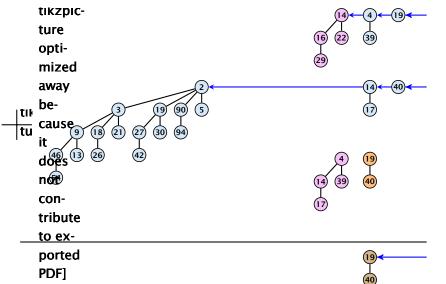


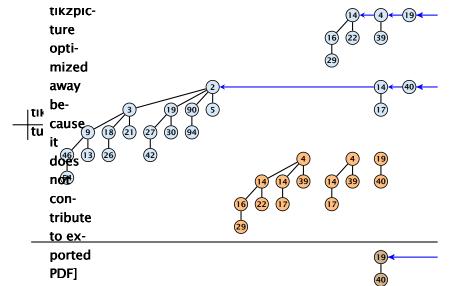


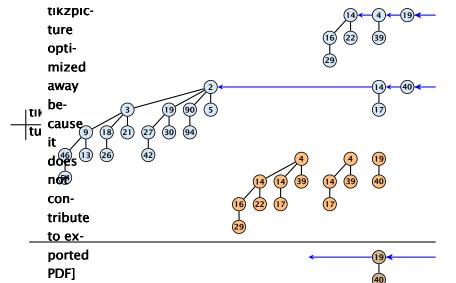


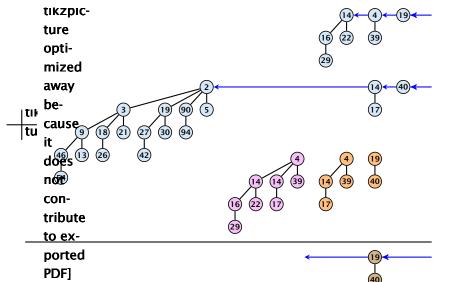


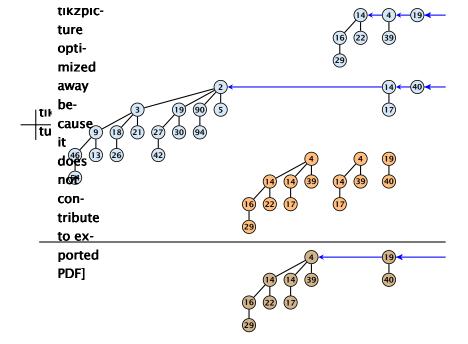


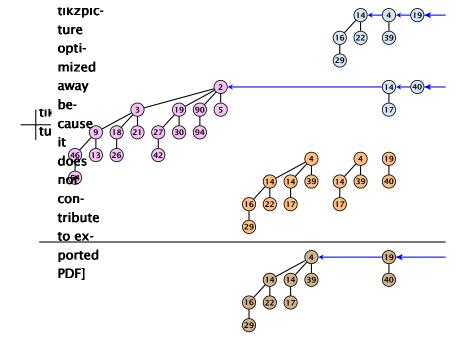


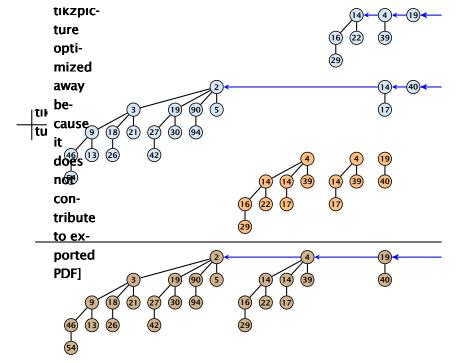


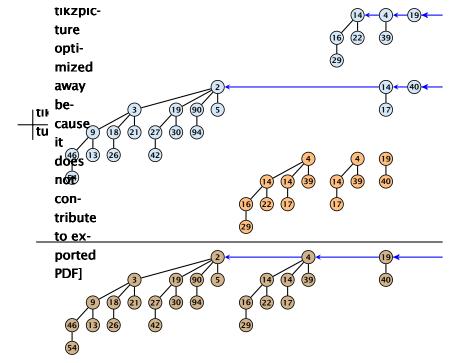












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• Create a new heap S' that contains just the element x.

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- Find the minimum key-value among all roots.
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- ▶ Time: $O(\log n)$ since the trees have height $O(\log n)$.

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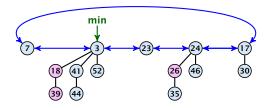
- ► Execute *S*. decrease-key(h, $-\infty$).
- ► Execute *S*. delete-min().

S. delete(handle *h*):

- **Execute** *S*. decrease-key $(h, -\infty)$.
- Execute *S*. delete-min().
- ▶ Time: $O(\log n)$.

Collection of trees that fulfill the heap property.

Structure is much more relaxed than binomial heaps.

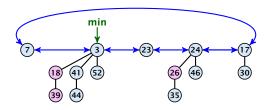


Additional implementation details:

- Every node x stores its degree in a field x. degree. Note that this can be updated in constant time when adding a child to x.
- Every node stores a boolean value x. marked that specifies whether x is marked or not.

The potential function:

- ightharpoonup t(S) denotes the number of trees in the heap.
- \blacktriangleright m(S) denotes the number of marked nodes.
- We use the potential function $\Phi(S) = t(S) + 2m(S)$.



The potential is $\Phi(S) = 5 + 2 \cdot 3 = 11$.

We assume that one unit of potential can pay for a constant amount of work, where the constant is chosen "big enough" (to take care of the constants that occur).

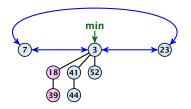
To make this more explicit we use c to denote the amount of work that a unit of potential can pay for.

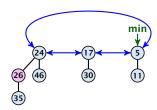
S. minimum()

- Access through the min-pointer.
- Actual cost $\mathcal{O}(1)$.
- No change in potential.
- ▶ Amortized cost $\mathcal{O}(1)$.

S. merge(S')

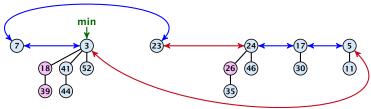
- Merge the root lists.
- Adjust the min-pointer





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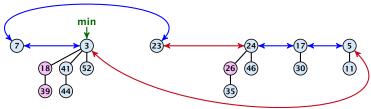


Running time:

Actual cost $\mathcal{O}(1)$.

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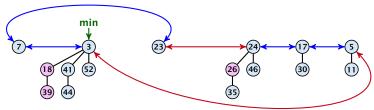


Running time:

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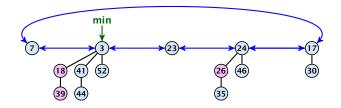
Running time:

- Actual cost $\mathcal{O}(1)$.
- No change in potential.
- \blacktriangleright Hence, amortized cost is $\mathcal{O}(1)$.



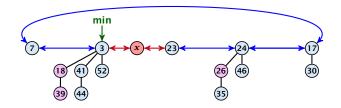
S.insert(x)

- ightharpoonup Create a new tree containing x.
- ► Insert *x* into the root-list.
- Update min-pointer, if necessary.



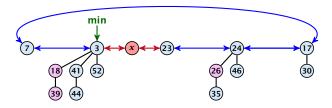
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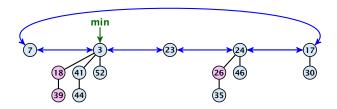
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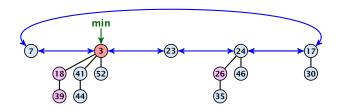
- Actual cost $\mathcal{O}(1)$.
- ightharpoonup Change in potential is +1.
- ▶ Amortized cost is c + O(1) = O(1).



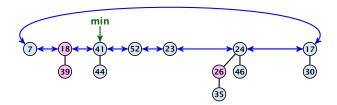


S. delete-min(x)

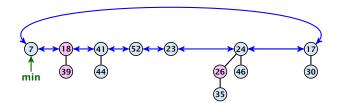
▶ Delete minimum; add child-trees to heap; time: $D(\min) \cdot \mathcal{O}(1)$.



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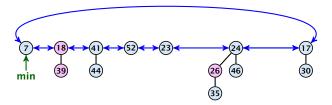


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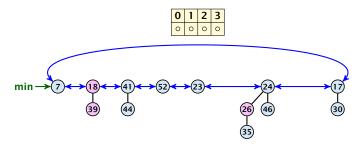


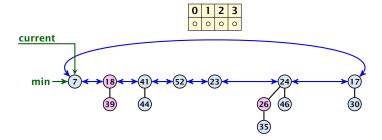
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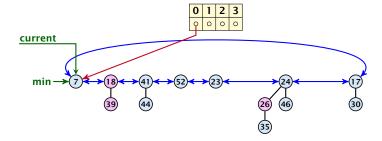
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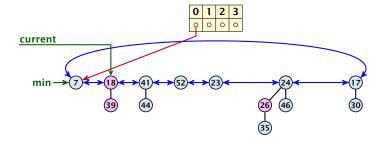


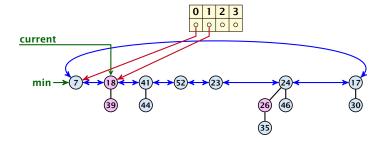
Consolidate root-list so that no roots have the same degree. Time $t\cdot\mathcal{O}(1)$ (see next slide).

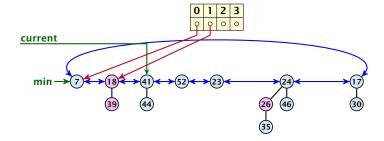


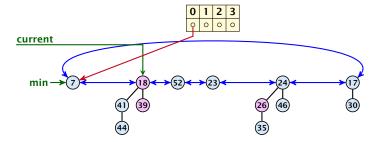


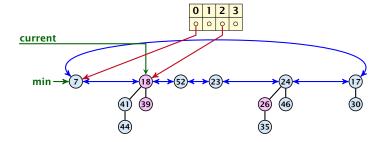


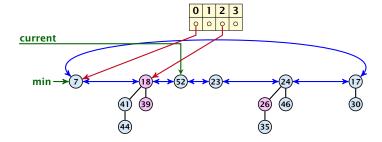


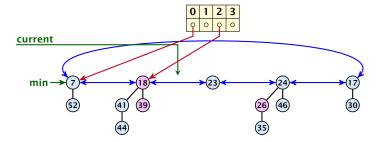


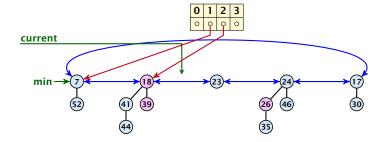


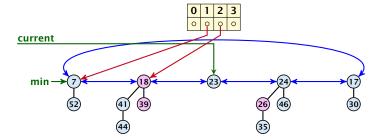


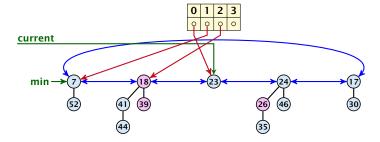


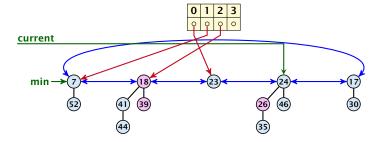


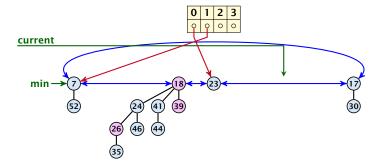


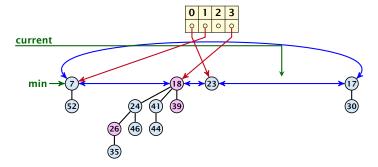


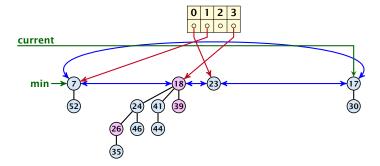


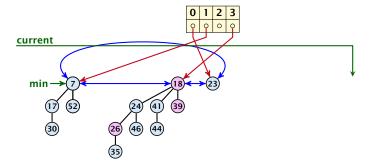


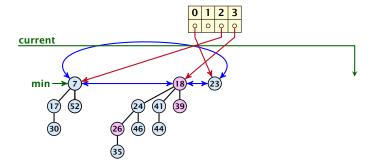


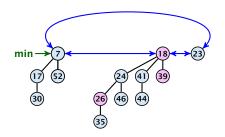












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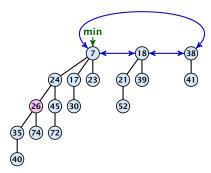
for $c \ge c_1$.



If the input trees of the consolidation procedure are binomial trees (for example only singleton vertices) then the output will be a set of distinct binomial trees, and, hence, the Fibonacci heap will be (more or less) a Binomial heap right after the consolidation.

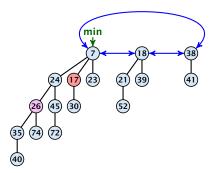
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If we do not have delete or decrease-key operations then $D_n \leq \log n$.



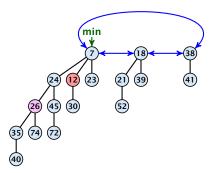
Case 1: decrease-key does not violate heap-property

Just decrease the key-value of element referenced by h. Nothing else to do.



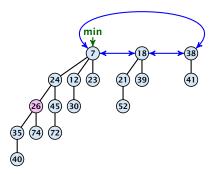
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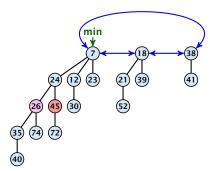
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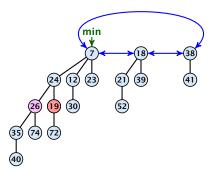
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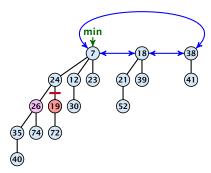
- Decrease key-value of element x reference by h.
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- Adjust min-pointers, if necessary.
- Mark the (previous) parent of x (unless it's a root).





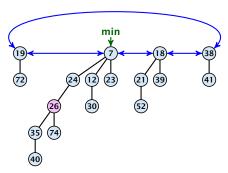
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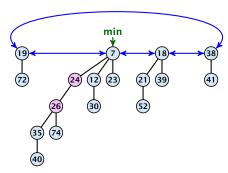
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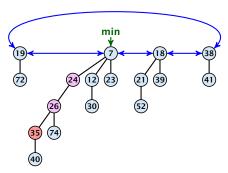
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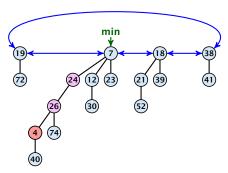
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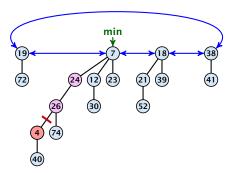
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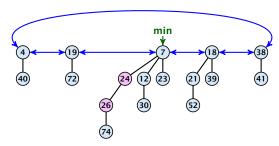
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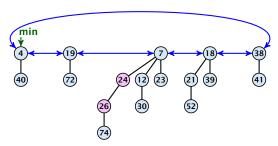
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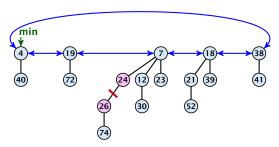
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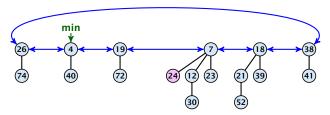
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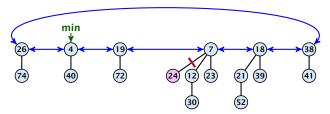
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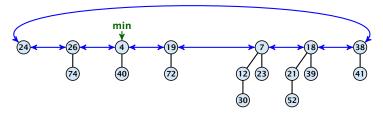
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- Execute the following:

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$$c_2(\ell+1)+c(4-\ell)$$

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- $t' = t + \ell$, as every cut creates one new root.
- ▶ $m' \le m (\ell 1) + 1 = m \ell + 2$, since all but the first cut unmarks a node; the last cut may mark a node.
- $\Delta \Phi \le \ell + 2(-\ell + 2) = 4 \ell$
- Amortized cost is at most

$$c_2(\ell+1) + c(4-\ell) \le (c_2-c)\ell + 4c + c_2$$

Actual cost:

- Constant cost for decreasing the value.
- ▶ Constant cost for each of ℓ cuts.
- ▶ Hence, cost is at most $c_2 \cdot (\ell + 1)$, for some constant c_2 .

- $ightharpoonup t'=t+\ell$, as every cut creates one new root.
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$$c_2(\ell+1) + c(4-\ell) \le (c_2-c)\ell + 4c + c_2 = \mathcal{O}(1)$$
, if $c \ge c_2$.

Delete node

H. delete(x):

- ▶ decrease value of x to $-\infty$.
- delete-min.

Amortized cost: $\mathcal{O}(D_n)$

- \triangleright $\mathcal{O}(1)$ for decrease-key.
- \triangleright $\mathcal{O}(D_n)$ for delete-min.

Lemma 21

Let x be a node with degree k and let y_1, \ldots, y_k denote the children of x in the order that they were linked to x. Then

$$degree(y_i) \ge \begin{cases} 0 & if i = 1\\ i - 2 & if i > 1 \end{cases}$$

Proof

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- Since, then y_i has lost at most one child.
- ▶ Therefore, degree(y_i) ≥ i 2.

Let s_k be the minimum possible size of a sub-tree rooted at a node of degree k that can occur in a Fibonacci heap.

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$$= 2 + \sum_{i=2}^{k-2} s_i$$

 $\phi=rac{1}{2}(1+\sqrt{5})$ denotes the *golden ratio*. Note that $\phi^2=1+\phi$.

Definition 22

Consider the following non-standard Fibonacci type sequence:

$$F_k = \begin{cases} 1 & \text{if } k = 0 \\ 2 & \text{if } k = 1 \\ F_{k-1} + F_{k-2} & \text{if } k \ge 2 \end{cases}$$

Facts:

- 1. $F_k \geq \phi^k$.
- **2.** For $k \ge 2$: $F_k = 2 + \sum_{i=0}^{k-2} F_i$.

The above facts can be easily proved by induction. From this it follows that $s_k \ge F_k \ge \phi^k$, which gives that the maximum degree in a Fibonacci heap is logarithmic.

k=0:
$$1 = F_0 \ge \Phi^0 = 1$$

k=1: $2 = F_1 \ge \Phi^1 \approx 1.61$
k-2,k-1 \rightarrow k: $F_k = F_{k-1} + F_{k-2} \ge \Phi^{k-1} + \Phi^{k-2} = \Phi^{k-2}(\Phi^{+1}) = \Phi^k$

k=2:
$$3 = F_2 = 2 + 1 = 2 + F_0$$

k-1 \rightarrow k: $F_k = F_{k-1} + F_{k-2} = 2 + \sum_{i=0}^{k-3} F_i + F_{k-2} = 2 + \sum_{i=0}^{k-2} F_i$



9 Union Find

Union Find Data Structure \mathcal{P} : Maintains a partition of disjoint sets over elements.

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Union Find Data Structure \mathcal{P} : Maintains a partition of disjoint sets over elements.

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- ▶ \mathcal{P} . find(x): Given a handle for an element x; find the set that contains x. Returns a representative/identifier for this set.
- ▶ \mathcal{P} . union(x, y): Given two elements x, and y that are currently in sets S_x and S_y , respectively, the function replaces S_x and S_y by $S_x \cup S_y$ and returns an identifier for the new set.

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Applications:

► Keep track of the connected components of a dynamic graph that changes due to insertion of nodes and edges.

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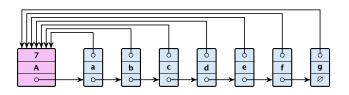
- Keep track of the connected components of a dynamic graph that changes due to insertion of nodes and edges.
- Kruskals Minimum Spanning Tree Algorithm

Algorithm 1 Kruskal-MST(G = (V, E), w)

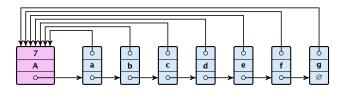
- 2: for all $v \in V$ do
- 3: $v. set \leftarrow P. makeset(v. label)$
- 4: sort edges in non-decreasing order of weight w
- 5: **for all** $(u, v) \in E$ in non-decreasing order **do**
- if \mathcal{P} . find(u. set) $\neq \mathcal{P}$. find(v. set) then
- $A \leftarrow A \cup \{(u,v)\}$
- \mathcal{P} . union(u. set, v. set)

► The elements of a set are stored in a list; each node has a backward pointer to the head.

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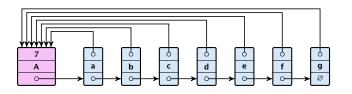


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- find(x) can be performed in constant time.

union(x, y)

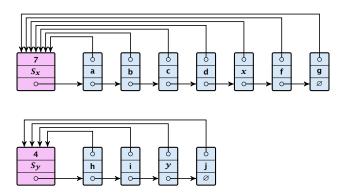
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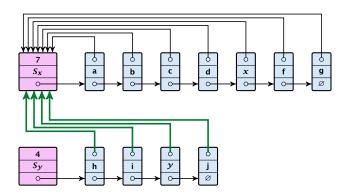
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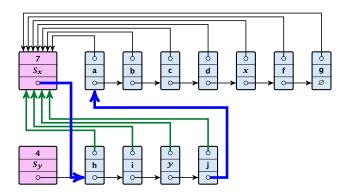
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- Adjust the size-field of list S_x .

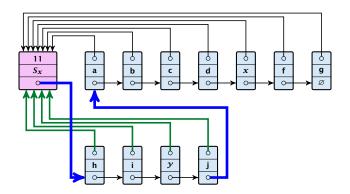
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- ▶ Insert list S_{γ} at the head of S_{χ} .
- Adjust the size-field of list S_x .
- ► Time: $\min\{|S_x|, |S_y|\}$.













Running times:

- ightharpoonup find(x): constant
- makeset(x): constant
- union(x, y): O(n), where n denotes the number of elements contained in the set system.

Lemma 23

The list implementation for the ADT union find fulfills the following amortized time bounds:

- find(x): $\mathcal{O}(1)$.
- ightharpoonup makeset(x): $O(\log n)$.
- union(x, y): O(1).

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- If we can find a charging scheme that guarantees that balances always stay positive the amortized time bounds are proven.

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- Later operations charge the account but the balance never drops below zero.

6. Feb. 2022

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- ▶ Charge c to every element in set S_x .



List Implementation

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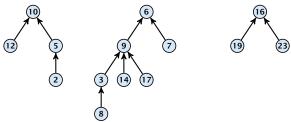
Proof.

Whenever an element x is charged the number of elements in x's set doubles. This can happen at most $|\log n|$ times.



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- The root of the tree is the label of the set.
- Only pointer to parent exists; we cannot list all elements of a given set.

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- Example:



Set system {2, 5, 10, 12}, {3, 6, 7, 8, 9, 14, 17}, {16, 19, 23}.

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Create a singleton tree. Return pointer to the root.

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- Start at element x in the tree. Go upwards until you reach the root.
- ► Time: $\mathcal{O}(\text{level}(x))$, where level(x) is the distance of element x to the root in its tree. Not constant.

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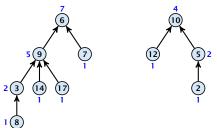
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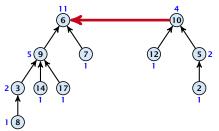
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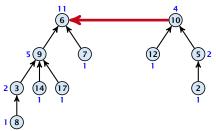
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▶ Time: constant for link(a, b) plus two find-operations.

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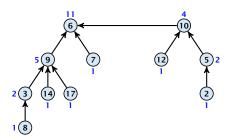
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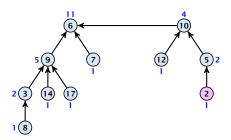
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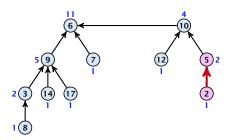
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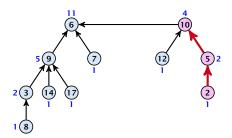
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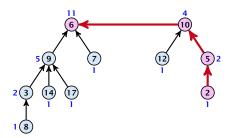
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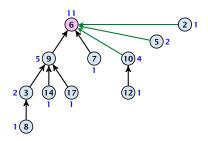
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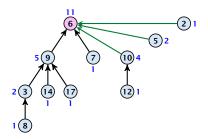


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Note that the size-fields now only give an upper bound on the size of a sub-tree.

Asymptotically the cost for a find-operation does not increase due to the path compression heuristic.

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However, for a worst-case analysis there is no improvement on the running time. It can still happen that a find-operation takes time $\mathcal{O}(\log n)$.

Amortized Analysis

Definitions:

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size(v) = the number of nodes that were in the sub-tree rooted at v when v became the child of another node (or the number of nodes if v is the root).

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Lemma 26

The rank of a parent must be strictly larger than the rank of a child.



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- This holds because the rank-sequence of the roots of the different trees that contain v during the running time of the algorithm is a strictly increasing sequence.



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- A node v sees at most one node of rank s during the running time of the algorithm.
- ▶ This holds because the rank-sequence of the roots of the different trees that contain v during the running time of the algorithm is a strictly increasing sequence.
- Hence, every node *sees* at most one rank s node, but every rank s node is seen by at least 2^s different nodes.

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We define

$$tow(i) := \left\{ \begin{array}{ll} 1 & \text{if } i = 0 \\ 2^{tow(i-1)} & \text{otw.} \end{array} \right.$$

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Theorem 28

Union find with path compression fulfills the following amortized running times:

- ightharpoonup makeset(x) : $\mathcal{O}(\log^*(n))$
- $ightharpoonup find(x) : \mathcal{O}(\log^*(n))$
- ightharpoonup union(x, y) : $\mathcal{O}(\log^*(n))$

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- ► The maximum non-empty rank group is $\log^*(\lfloor \log n \rfloor) \leq \log^*(n) 1$ (which holds for $n \geq 2$).
- ▶ Hence, the total number of rank-groups is at most $\log^* n$.

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- Otherwise we charge the cost to the find-account.

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- After a node v is charged its parent-edge is re-assigned. The rank of the parent strictly increases.
- After some charges to v the parent will be in a larger rank-group. $\Rightarrow v$ will never be charged again.
- ► The total charge made to a node in rank-group g is at most $tow(g) tow(g-1) 1 \le tow(g)$.

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The total charge is at most

$$\sum_{g} n(g) \cdot \text{tow}(g) ,$$

where n(g) is the number of nodes in group g.

For $g \ge 1$ we have

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$$n(g) \le \sum_{s=\text{tow}(g-1)+1}^{\text{tow}(g)} \frac{n}{2^s}$$

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Hence,

$$\sum_{g} n(g) \operatorname{tow}(g) \le n(0) \operatorname{tow}(0) + \sum_{g>1} n(g) \operatorname{tow}(g) \le n \log^*(n)$$

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This means if we inflate the cost of makeset to $\log^* n$ and add this to the node account of v then the balances of all node accounts will sum up to a positive value (this is sufficient to obtain an amortized bound).



The analysis is not tight. In fact it has been shown that the amortized time for the union-find data structure with path compression is $\mathcal{O}(\alpha(m,n))$, where $\alpha(m,n)$ is the inverse Ackermann function which grows a lot lot slower than $\log^* n$. (Here, we consider the average running time of m operations on at most n elements).

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There is also a lower bound of $\Omega(\alpha(m, n))$.

$$A(x,y) = \begin{cases} y+1 & \text{if } x = 0 \\ A(x-1,1) & \text{if } y = 0 \\ A(x-1,A(x,y-1)) & \text{otw.} \end{cases}$$

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$$\alpha(m,n) = \min\{i \ge 1 : A(i,\lfloor m/n \rfloor) \ge \log n\}$$

- A(0, v) = v + 1
- A(1, v) = v + 2
- $A(2, \nu) = 2\nu + 3$
- ► $A(3, y) = 2^{y+3} 3$ ► $A(4, y) = 2^{2^{2^2}} 3$