WS 2024/25

Efficient Algorithms

Harald Räcke

Fakultät für Informatik TU München

https://www.moodle.tum.de/course/view.php?id=100478

Winter Term 2024/25

Part I

Organizational Matters

Part I

Organizational Matters

Modul: IN2003

Name: "Efficient Algorithms and Data Structures" "Effiziente Algorithmen und Datenstrukturen"

ECTS: 8 Credit points

Lectures:

4 SWS
 Mon 10:00-12:00 (Room Interim2)
 Fri 10:00-12:00 (Room Interim2)

Webpage: https://www.moodle.tum.de/course/view.php?id=100478

- Required knowledge:
 - ► IN0001, IN0003
 - "Introduction to Informatics 1/2"
 - "Einführung in die Informatik 1/2"
 - ► IN0007
 - "Fundamentals of Algorithms and Data Structures"
 - "Grundlagen: Algorithmen und Datenstrukturen" (GAD)
 - ► IN0011
 - "Basic Theoretic Informatics"
 - "Einführung in die Theoretische Informatik" (THEO)
 - ► IN0015
 - "Discrete Structures"
 - "Diskrete Strukturen" (DS)
 - ► IN0018
 - "Discrete Probability Theory"
 - "Diskrete Wahrscheinlichkeitstheorie" (DWT)



12. Jan. 2025

The Lecturer

- ► Harald Räcke
- Email: raecke@in.tum.de
- Room: 03.09.044
- Office hours: (by appointment)

Tutorials

- Omar AbdelWanis
- omar.abdelwanis@in.tum.de
- Room: 03.09.042
- Office hours: (by appointment)

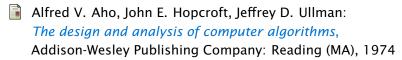


1 Contents

- Foundations
 - Machine models
 - Efficiency measures
 - Asymptotic notation
 - Recursion
- Higher Data Structures
 - Search trees
 - Hashing
 - Priority queues
 - Union/Find data structures
- Cuts/Flows
- Matchings



2 Literatur



Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein:

Introduction to algorithms,

McCraw-Hill 1990

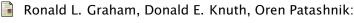
McGraw-Hill, 1990

Michael T. Goodrich, Roberto Tamassia:

Algorithm design: Foundations, analysis, and internet examples,
John Wiley & Sons, 2002



2 Literatur



Concrete Mathematics,

2. Auflage, Addison-Wesley, 1994

Volker Heun:

Grundlegende Algorithmen: Einführung in den Entwurf und die Analyse effizienter Algorithmen,

2. Auflage, Vieweg, 2003

Jon Kleinberg, Eva Tardos:

Algorithm Design,

Addison-Wesley, 2005

Donald E. Knuth:

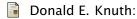
The art of computer programming. Vol. 1: Fundamental Algorithms,

3. Auflage, Addison-Wesley, 1997



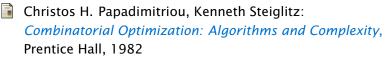
12. Jan. 2025

2 Literatur



The art of computer programming. Vol. 3: Sorting and Searching,

3. Auflage, Addison-Wesley, 1997



Uwe Schöning: Algorithmik, Spektrum Akademischer Verlag, 2001

Steven S. Skiena:
The Algorithm Design Manual,
Springer, 1998

Part II

Foundations

3 Goals

- Gain knowledge about efficient algorithms for important problems, i.e., learn how to solve certain types of problems efficiently.
- Learn how to analyze and judge the efficiency of algorithms.
- Learn how to design efficient algorithms.

12. Jan. 2025

12/530

What do you measure?

- Memory requirement
- Running time
- Number of comparisons
- Number of multiplications
- Number of hard-disc accesses
- Program size
- Power consumption
- **.**..



How do you measure?

- Implementing and testing on representative inputs
 - How do you choose your inputs?
 - May be very time-consuming.
 - Very reliable results if done correctly.
 - Results only hold for a specific machine and for a specific set of inputs.
- Theoretical analysis in a specific model of computation.
 - Gives asymptotic bounds like "this algorithm always runs in time $\mathcal{O}(n^2)$ ".
 - Typically focuses on the worst case.
 - Can give lower bounds like "any comparison-based sorting algorithm needs at least $\Omega(n \log n)$ comparisons in the worst case".



Input length

The theoretical bounds are usually given by a function $f: \mathbb{N} \to \mathbb{N}$ that maps the input length to the running time (or storage space, comparisons, multiplications, program size etc.).

The input length may e.g. be

- the size of the input (number of bits)
- the number of arguments

Example 1

Suppose n numbers from the interval $\{1,\ldots,N\}$ have to be sorted. In this case we usually say that the input length is n instead of e.g. $n\log N$, which would be the number of bits required to encode the input.



Model of Computation

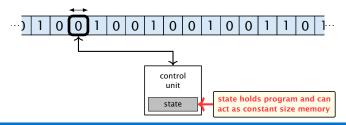
How to measure performance

- Calculate running time and storage space etc. on a simplified, idealized model of computation, e.g. Random Access Machine (RAM), Turing Machine (TM), . . .
- 2. Calculate number of certain basic operations: comparisons, multiplications, harddisc accesses, . . .

Version 2. is often easier, but focusing on one type of operation makes it more difficult to obtain meaningful results.

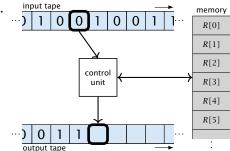
Turing Machine

- Very simple model of computation.
- Only the "current" memory location can be altered.
- Very good model for discussing computability, or polynomial vs. exponential time.
- Some simple problems like recognizing whether input is of the form xx, where x is a string, have quadratic lower bound.
- \Rightarrow Not a good model for developing efficient algorithms.



Random Access Machine (RAM)

- Input tape and output tape (sequences of zeros and ones; unbounded length).
- Memory unit: infinite but countable number of registers $R[0], R[1], R[2], \ldots$
- Registers hold integers.
- Indirect addressing.



Note that in the picture on the right the tapes are one-directional, and that a READ- or WRITE-operation always ad-

Random Access Machine (RAM)

Operations

- ▶ input operations (input tape $\rightarrow R[i]$)
 - ► READ *i*
- output operations $(R[i] \rightarrow \text{output tape})$
 - ► WRITE *i*
- register-register transfers
 - ightharpoonup R[j] := R[i]
 - ightharpoonup R[j] := 4
- indirect addressing
 - ▶ R[j] := R[R[i]] loads the content of the R[i]-th register into the j-th register
 - R[R[i]] := R[j] loads the content of the j-th into the R[i]-th register

Random Access Machine (RAM)

Operations

- branching (including loops) based on comparisons
 - jump x jumps to position x in the program; sets instruction counter to x; reads the next operation to perform from register R[x]
 - jumpz x R[i]
 jump to x if R[i] = 0
 - if not the instruction counter is increased by 1;
 - jumpi i
 jump to R[i] (indirect jump);
- arithmetic instructions: +, -, ×, /
 - ► R[i] := R[j] + R[k];R[i] := -R[k];

The jump-directives are very close to the jump-instructions contained in the assembler language of real machines.

20/530

Model of Computation

- uniform cost modelEvery operation takes time 1.
- logarithmic cost model The cost depends on the content of memory cells:
 - The time for a step is equal to the largest operand involved;
 - ► The storage space of a register is equal to the length (in bits) of the largest value ever stored in it.

Bounded word RAM model: cost is uniform but the largest value stored in a register may not exceed 2^w , where usually $w = \log_2 n$.

The latter model is quite realistic as the word-size of a standard computer that handles a problem of size n must be at least $\log_2 n$ as otherwise the computer could either not store the problem instance or not address all its memory.

Example 2

Algorithm 1 RepeatedSquaring(n)

1: $r \leftarrow 2$; 2: **for** $i = 1 \rightarrow n$ **do** 3: $r \leftarrow r^2$ 4: **return** r

- running time (for Line 3):
 - uniform model: n steps
 - logarithmic model:

$$2+3+5+\cdots+(1+2^n)=2^{n+1}-1+n=\Theta(2^n)$$

- space requirement:
 - uniform model: $\mathcal{O}(1)$
 - logarithmic model: $\mathcal{O}(2^n)$



There are different types of complexity bounds:

best-case complexity:

$$C_{\mathrm{bc}}(n) := \min\{C(x) \mid |x| = n\}$$

Usually easy to analyze, but not very meaningful.

worst-case complexity:

$$C_{\mathrm{WC}}(n) := \max\{C(x) \mid |x| = n\}$$

Usually moderately easy to analyze; sometimes too pessimistic.

average case complexity:

$$C_{\text{avg}}(n) := \frac{1}{|I_n|} \sum_{|x|=n} C(x)$$

more general: probability measure μ

$$\mu$$
 is a probability distribution over inputs of length n . $C_{\mathrm{avg}}(n) := \sum_{x \in I_n} \mu(x) \cdot C(x)$

 $C(x) \begin{cases} \cos t \text{ of instance} \\ x \end{cases}$ $|x| \quad \text{input length of instance } x$

 I_n set of instances of length n

23/530

There are different types of complexity bounds:

- amortized complexity:
 - The average cost of data structure operations over a worst case sequence of operations.
- randomized complexity:

The algorithm may use random bits. Expected running time (over all possible choices of random bits) for a fixed input x.

Then take the worst-case over all x with |x| = n.

	C(x)	cost of instance
		X
	<i>x</i>	input length of
		instance x
	I_n	set of instances
		of length n

Bibliography

[MS08] Kurt Mehlhorn, Peter Sanders:

Algorithms and Data Structures — The Basic Toolbox, Springer, 2008

[CLRS90] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein: Introduction to algorithms (3rd ed.),

McGraw-Hill. 2009

Chapter 2.1 and 2.2 of [MS08] and Chapter 2 of [CLRS90] are relevant for this section.

We are usually not interested in exact running times, but only in an asymptotic classification of the running time, that ignores constant factors and constant additive offsets.

- We are usually interested in the running times for large values of n. Then constant additive terms do not play an important role.
- An exact analysis (e.g. exactly counting the number of operations in a RAM) may be hard, but wouldn't lead to more precise results as the computational model is already quite a distance from reality.
- A linear speed-up (i.e., by a constant factor) is always possible by e.g. implementing the algorithm on a faster machine.
- Running time should be expressed by simple functions.



Formal Definition

Let f, g denote functions from \mathbb{N} to \mathbb{R}^+ .

- ▶ $\mathcal{O}(f) = \{g \mid \exists c > 0 \ \exists n_0 \in \mathbb{N}_0 \ \forall n \geq n_0 : [g(n) \leq c \cdot f(n)]\}$ (set of functions that asymptotically grow not faster than f)
- ▶ $\Omega(f) = \{g \mid \exists c > 0 \ \exists n_0 \in \mathbb{N}_0 \ \forall n \geq n_0 : [g(n) \geq c \cdot f(n)]\}$ (set of functions that asymptotically grow not slower than f)
- $\Theta(f) = \Omega(f) \cap \mathcal{O}(f)$ (functions that asymptotically have the same growth as f)
- ▶ $o(f) = \{g \mid \forall c > 0 \ \exists n_0 \in \mathbb{N}_0 \ \forall n \geq n_0 : [g(n) \leq c \cdot f(n)]\}$ (set of functions that asymptotically grow slower than f)
- ▶ $\omega(f) = \{g \mid \forall c > 0 \ \exists n_0 \in \mathbb{N}_0 \ \forall n \geq n_0 : [g(n) \geq c \cdot f(n)]\}$ (set of functions that asymptotically grow faster than f)

25/530

There is an equivalent definition using limes notation (assuming that the respective limes exists). f and g are functions from \mathbb{N}_0 to \mathbb{R}_0^+ .

$$ightharpoonup g \in \mathcal{O}(f): \quad 0 \le \lim_{n \to \infty} \frac{g(n)}{f(n)} < \infty$$

•
$$g \in \Omega(f)$$
: $0 < \lim_{n \to \infty} \frac{g(n)}{f(n)} \le \infty$

$$g \in o(f): \quad \lim_{n \to \infty} \frac{g(n)}{f(n)} = 0$$

•
$$g \in \omega(f)$$
: $\lim_{n \to \infty} \frac{g(n)}{f(n)} = \infty$

- Note that for the version of the Landau notation defined here, we assume that f and g are positive functions.
- There also exist versions for arbitrary functions, and for the case that the limes is not infinity.

Abuse of notation

- 1. People write $f = \mathcal{O}(g)$, when they mean $f \in \mathcal{O}(g)$. This is **not** an equality (how could a function be equal to a set of functions).
- **2.** People write $f(n) = \mathcal{O}(g(n))$, when they mean $f \in \mathcal{O}(g)$, with $f: \mathbb{N} \to \mathbb{R}^+$, $n \mapsto f(n)$, and $g: \mathbb{N} \to \mathbb{R}^+$, $n \mapsto g(n)$.
- 3. People write e.g. h(n) = f(n) + o(g(n)) when they mean that there exists a function $z: \mathbb{N} \to \mathbb{R}^+, n \mapsto z(n), z \in o(g)$ such that h(n) = f(n) + z(n).

- **2.** In this context f(n) does **not** mean the function f evaluated at n, but instead it is a shorthand for the function itself (leaving out do-1
 - to ignore constant factors. For example the median of n elements can be determined main and codomain and only giving the rule using $\frac{3}{2}n + o(n)$ comparisons. of correspondence of the function).

3. This is particularly useful if you do not want

Abuse of notation

4. People write $\mathcal{O}(f(n)) = \mathcal{O}(g(n))$, when they mean $\mathcal{O}(f(n)) \subseteq \mathcal{O}(g(n))$. Again this is not an equality.

In this context f(n) does not mean the function f evaluated at n, but instead it is a shorthand for the function itself (leaving out do-

main and codomain and only giving the rule

of correspondence of the function).

3. This is particularly useful if you do not want to ignore constant factors. For example the median of n elements can be determined using $\frac{3}{2}n + o(n)$ comparisons.

How do we interpret an expression like:

$$2n^2 + 3n + 1 = 2n^2 + \Theta(n)$$

Here, $\Theta(n)$ stands for an anonymous function in the set $\Theta(n)$ that makes the expression true.

Note that $\Theta(n)$ is on the right hand side, otw. this interpretation is wrong.

How do we interpret an expression like:

$$2n^2 + \mathcal{O}(n) = \Theta(n^2)$$

Regardless of how we choose the anonymous function $f(n) \in \mathcal{O}(n)$ there is an anonymous function $g(n) \in \Theta(n^2)$ that makes the expression true.

The $\Theta(i)$ -symbol on the left represents one anonymous function $f: \mathbb{N} \to \mathbb{R}^+$, and then $\sum_i f(i)$ is computed.

How do we interpret an expression like:

$$\sum_{i=1}^n \Theta(i) = \Theta(n^2)$$

Careful!

"It is understood" that every occurrence of an \mathcal{O} -symbol (or $\Theta, \Omega, o, \omega$) on the left represents one anonymous function.

Hence, the left side is not equal to

$$\Theta(1) + \Theta(2) + \cdots + \Theta(n-1) + \Theta(n)$$

 $\Theta(1) + \Theta(2) + \cdots + \Theta(n-1) + \Theta(n)$ does not really have a reasonable interpreta-

We can view an expression containing asymptotic notation as generating a set:

$$n^2 \cdot \mathcal{O}(n) + \mathcal{O}(\log n)$$

represents

$$\begin{split} \left\{f: \mathbb{N} \to \mathbb{R}^+ \mid f(n) = n^2 \cdot g(n) + h(n) \\ & \text{with } g(n) \in \mathcal{O}(n) \text{ and } h(n) \in \mathcal{O}(\log n) \right\} \\ & \text{Recall that according to the previous} \\ & \text{slide e.g. the expressions } \sum_{i=1}^n \mathcal{O}(i) \text{ and} \\ & \sum_{i=1}^{n/2} \mathcal{O}(i) + \sum_{i=n/2+1}^n \mathcal{O}(i) \text{ generate difference} \end{split}$$

Then an asymptotic equation can be interpreted as containement btw. two sets:

$$n^2 \cdot \mathcal{O}(n) + \mathcal{O}(\log n) = \Theta(n^2)$$

represents

$$n^2 \cdot \mathcal{O}(n) + \mathcal{O}(\log n) \subseteq \Theta(n^2)$$

Note that the equation does not hold.

Lemma 3

Let f, g be functions with the property

 $\exists n_0 > 0 \ \forall n \ge n_0 : f(n) > 0$ (the same for g). Then

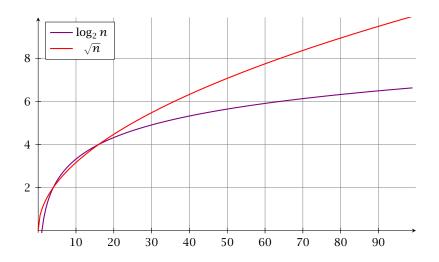
- $c \cdot f(n) \in \Theta(f(n))$ for any constant c
- $\bullet \ \mathcal{O}(f(n)) \cdot \mathcal{O}(g(n)) = \mathcal{O}(f(n) \cdot g(n))$
- $\mathcal{O}(f(n)) + \mathcal{O}(g(n)) = \mathcal{O}(\max\{f(n), g(n)\})$

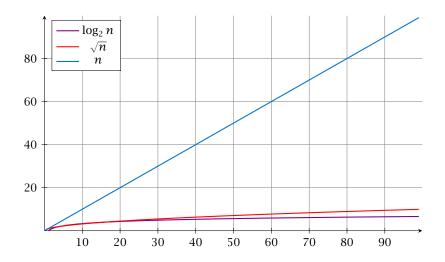
The expressions also hold for Ω . Note that this means that $f(n) + g(n) \in \Theta(\max\{f(n), g(n)\})$.

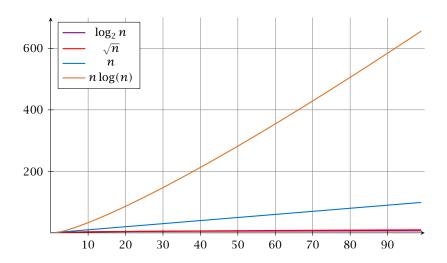
Asymptotic Notation

Comments

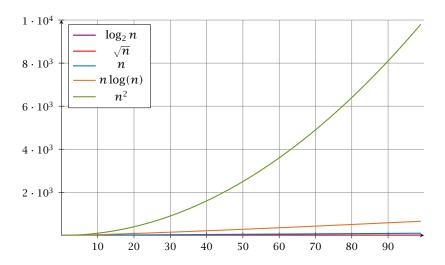
- Do not use asymptotic notation within induction proofs.
- For any constants a, b we have $\log_a n = \Theta(\log_b n)$. Therefore, we will usually ignore the base of a logarithm within asymptotic notation.
- In general $\log n = \log_2 n$, i.e., we use 2 as the default base for the logarithm.



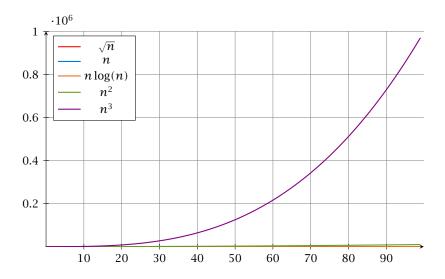


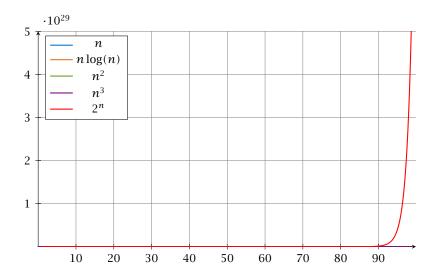












Laufzeiten

Funktion	Eingabelänge n								
f(n)	10	10^{2}	10^{3}	10^{4}	104 105		10 ⁷	108	
$\log n$	33 ns	66ns	0.1µs	0.1µs	0.2µs	0.2µs	0.2µs	0.3µs	
\sqrt{n}	32ns	$0.1 \mu s$	0.3µs	1µs	$3.1 \mu s$	10µs	31 µs	$0.1 \mathrm{ms}$	
n	100ns	1µs	10 µs	$0.1 \mathrm{ms}$	1ms	10ms	0.1s	1s	
$n \log n$	0.3µs	6.6µs	0.1ms	1.3ms	16ms	0.2s	2.3s	27s	
$n^{3/2}$	0.3µs	10µs	0.3ms	10ms	0.3s	10s	5.2min	2.7h	
n^2	1µs	$0.1 \mathrm{ms}$	10ms	1s	1.7min	2.8h	11 d	3.2 y	
n^3	10µs	10ms	10s	2.8h	115 d	317 y	3.2·10 ⁵ y		
1.1^{n}	26ns	0.1 ms	$7.8 \cdot 10^{25}$ y						
2^n	10µs	$4\cdot 10^{14}$ y							
n!	36ms	$3 \cdot 10^{142}$ y							

1 Operation = 10ns; 100MHz

Alter des Universums: ca. $13.8 \cdot 10^9 \mathrm{y}$

Asymptotic Notation

In general asymptotic classification of running times is a good measure for comparing algorithms:

- ▶ If the running time analysis is tight and actually occurs in practise (i.e., the asymptotic bound is not a purely theoretical worst-case bound), then the algorithm that has better asymptotic running time will always outperform a weaker algorithm for large enough values of *n*.
- However, suppose that I have two algorithms:
 - Algorithm A. Running time $f(n) = 1000 \log n = \mathcal{O}(\log n)$.
 - ► Algorithm B. Running time $g(n) = \log^2 n$.

Clearly f = o(g). However, as long as $\log n \le 1000$ Algorithm B will be more efficient.



Multiple Variables in Asymptotic Notation

Sometimes the input for an algorithm consists of several parameters (e.g., nodes and edges of a graph (n and m)).

If we want to make asymptoic statements for $n \to \infty$ and $m \to \infty$ we have to extend the definition to multiple variables.

Formal Definition

Let f, g denote functions from \mathbb{N}^d to \mathbb{R}_0^+ .

 $\mathcal{O}(f) = \{g \mid \exists c > 0 \ \exists N \in \mathbb{N}_0 \ \forall \vec{n} \ \text{with} \ n_i \geq N \ \text{for some} \ i : \\ [g(\vec{n}) \leq c \cdot f(\vec{n})] \}$

(set of functions that asymptotically grow not faster than f)

Multiple Variables in Asymptotic Notation

Example 4

- ▶ $f: \mathbb{N} \to \mathbb{R}_0^+$, f(n,m) = 1 und $g: \mathbb{N} \to \mathbb{R}_0^+$, g(n,m) = n-1 then $f = \mathcal{O}(g)$ does not hold
- ▶ $f: \mathbb{N} \to \mathbb{R}_0^+$, f(n,m) = 1 und $g: \mathbb{N} \to \mathbb{R}_0^+$, g(n,m) = n then: $f = \mathcal{O}(g)$
- ▶ $f: \mathbb{N}_0 \to \mathbb{R}_0^+$, f(n,m) = 1 und $g: \mathbb{N}_0 \to \mathbb{R}_0^+$, g(n,m) = n then $f = \mathcal{O}(g)$ does not hold

5 Asymptotic Notation

Bibliography

[MS08] Kurt Mehlhorn, Peter Sanders:

Algorithms and Data Structures — The Basic Toolbox, Springer, 2008

[CLRS90] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein:

Introduction to algorithms (3rd ed.),

McGraw-Hill, 2009

Mainly Chapter 3 of [CLRS90]. [MS08] covers this topic in chapter 2.1 but not very detailed.

6 Recurrences

Algorithm 2 mergesort(list *L*)

1: $n \leftarrow \text{size}(L)$ 2: **if** $n \le 1$ **return** L3: $L_1 \leftarrow L[1 \cdots \lfloor \frac{n}{2} \rfloor]$ 4: $L_2 \leftarrow L[\lfloor \frac{n}{2} \rfloor + 1 \cdots n]$ 5: mergesort(L_1)

6: mergesort(L_2)
7: $L \leftarrow \text{merge}(L_1, L_2)$ 8: **return** L

This algorithm requires

$$T(n) = T\left(\left\lceil \frac{n}{2}\right\rceil\right) + T\left(\left\lfloor \frac{n}{2}\right\rfloor\right) + \mathcal{O}(n) \le 2T\left(\left\lceil \frac{n}{2}\right\rceil\right) + \mathcal{O}(n)$$

comparisons when n > 1 and 0 comparisons when $n \le 1$.

Recurrences

How do we bring the expression for the number of comparisons (≈ running time) into a closed form?

For this we need to solve the recurrence.

Methods for Solving Recurrences

1. Guessing+Induction

Guess the right solution and prove that it is correct via induction. It needs experience to make the right guess.

2. Master Theorem

For a lot of recurrences that appear in the analysis of algorithms this theorem can be used to obtain tight asymptotic bounds. It does not provide exact solutions.

3. Characteristic Polynomial

Linear homogenous recurrences can be solved via this method.



Methods for Solving Recurrences

4. Generating Functions

A more general technique that allows to solve certain types of linear inhomogenous relations and also sometimes non-linear recurrence relations.

5. Transformation of the Recurrence

Sometimes one can transform the given recurrence relations so that it e.g. becomes linear and can therefore be solved with one of the other techniques.



First we need to get rid of the \mathcal{O} -notation in our recurrence:

$$T(n) \le \begin{cases} 2T(\lceil \frac{n}{2} \rceil) + cn & n \ge 2\\ 0 & \text{otherwise} \end{cases}$$

Informal way:

Assume that instead we have

$$T(n) \le \begin{cases} 2T(\frac{n}{2}) + cn & n \ge 2\\ 0 & \text{otherwise} \end{cases}$$

One way of solving such a recurrence is to guess a solution, and check that it is correct by plugging it in.

Suppose we guess $T(n) \le dn \log n$ for a constant d. Then

$$T(n) \le 2T\left(\frac{n}{2}\right) + cn$$

$$\le 2\left(d\frac{n}{2}\log\frac{n}{2}\right) + cn$$

$$= dn(\log n - 1) + cn$$

$$= dn\log n + (c - d)n$$

$$\le dn\log n$$

if we choose d > c.

Formally, this is not correct if n is not a power of 2. Also even in this case one would need to do an induction proof.

$$T(n) \le \begin{cases} 2T(\frac{n}{2}) + cn & n \ge 16 \\ b & \text{otw.} \end{cases}$$

Guess: $T(n) \le dn \log n$.

Proof. (by induction)

- **base case** $(2 \le n < 16)$: true if we choose $d \ge b$.
- ▶ induction step $n/2 \rightarrow n$:

Let $n = 2^k \ge 16$. Suppose statem. is true for n' = n/2. We prove it for n:

$$T(n) \le 2T\left(\frac{n}{2}\right) + cn$$

$$\le 2\left(d\frac{n}{2}\log\frac{n}{2}\right) + cn$$

$$= dn(\log n - 1) + cn$$

$$= dn\log n + (c - d)n$$

$$\le dn\log n$$

Note that this proves the statement for n = 2^k, k ∈ N≥1, as the statement is wrong for n = 1.
The base case is usually omitted, as it is the same for different recurrences.

Hence, statement is true if we choose $d \ge c$.

How do we get a result for all values of n?

We consider the following recurrence instead of the original one:

$$T(n) \le \begin{cases} 2T(\lceil \frac{n}{2} \rceil) + cn & n \ge 16 \\ b & \text{otherwise} \end{cases}$$

Note that we can do this as for constant-sized inputs the running time is always some constant (b in the above case).

We also make a guess of $T(n) \le dn \log n$ and get

$$T(n) \leq 2T\left(\left\lceil\frac{n}{2}\right\rceil\right) + cn$$

$$\leq 2\left(d\left\lceil\frac{n}{2}\right\rceil\log\left\lceil\frac{n}{2}\right\rceil\right) + cn$$

$$\left\lceil\frac{n}{2}\right\rceil \leq \frac{n}{2} + 1\right\rceil \leq 2\left(d(n/2 + 1)\log(n/2 + 1)\right) + cn$$

$$\left\lceil\frac{n}{2} + 1 \leq \frac{9}{16}n\right\rceil \leq dn\log\left(\frac{9}{16}n\right) + 2d\log n + cn$$

$$\left\lceil\log\frac{9}{16}n\right\rceil = \log n + (\log 9 - 4)\right\rceil = dn\log n + (\log 9 - 4)dn + 2d\log n + cn$$

$$\left\lceil\log n \leq \frac{n}{4}\right\rceil \leq dn\log n + (\log 9 - 3.5)dn + cn$$

$$\leq dn\log n - 0.33dn + cn$$

$$\leq dn\log n$$

for a suitable choice of d.

Lemma 5

Let $a \ge 1, b > 1$ and $\epsilon > 0$ denote constants. Consider the recurrence

$$T(n) = aT\left(\frac{n}{h}\right) + f(n)$$
.

Case 1.

If
$$f(n) = \mathcal{O}(n^{\log_b(a) - \epsilon})$$
 then $T(n) = \Theta(n^{\log_b a})$.

Case 2.

If
$$f(n) = \Theta(n^{\log_b(a)} \log^k n)$$
 then $T(n) = \Theta(n^{\log_b a} \log^{k+1} n)$, $k \ge 0$.

Case 3.

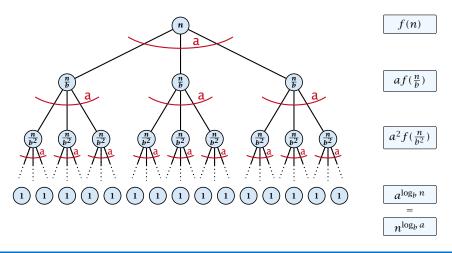
If
$$f(n) = \Omega(n^{\log_b(a) + \epsilon})$$
 and for sufficiently large n $af(\frac{n}{b}) \le cf(n)$ for some constant $c < 1$ then $T(n) = \Theta(f(n))$.

6.2 Master Theorem

We prove the Master Theorem for the case that n is of the form b^{ℓ} , and we assume that the non-recursive case occurs for problem size 1 and incurs cost 1.

The Recursion Tree

The running time of a recursive algorithm can be visualized by a recursion tree:



6.2 Master Theorem

This gives

$$T(n) = n^{\log_b a} + \sum_{i=0}^{\log_b n-1} a^i f\left(\frac{n}{b^i}\right) .$$

Case 1. Now suppose that $f(n) \le c n^{\log_b a - \epsilon}$.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$\log_b n - 1$$

$$\frac{b^{-i(\log_b a - \epsilon)} = b^{\epsilon i}(b^{\log_b a})^{-i} = b^{\epsilon i}a^{-i}}{\sum_{i=0}^{k} q^i = \frac{q^{k+1} - 1}{q - 1}} = cn^{\log_b a - \epsilon} \sum_{i=0}^{\log_b n - 1} (b^{\epsilon})^i$$

$$= cn^{\log_b a - \epsilon} (b^{\epsilon \log_b n} - 1)/(b^{\epsilon} - 1)$$

$$= cn^{\log_b a - \epsilon} (n^{\epsilon} - 1)/(b^{\epsilon} - 1)$$

$$= \frac{c}{b^{\epsilon} - 1} n^{\log_b a} (n^{\epsilon} - 1)/(n^{\epsilon})$$

Hence,

$$T(n) \le \left(\frac{c}{h^{\epsilon} - 1} + 1\right) n^{\log_b(a)}$$

$$\Rightarrow T(n) = \mathcal{O}(n^{\log_b a}).$$

Case 2. Now suppose that $f(n) \leq c n^{\log_b a}$.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$= c n^{\log_b a} \log_b n$$

Hence,

$$T(n) = \mathcal{O}(n^{\log_b a} \log_b n)$$

$$T(n) = \mathcal{O}(n^{\log_b a} \log_b n)$$
 $\Rightarrow T(n) = \mathcal{O}(n^{\log_b a} \log n).$

Case 2. Now suppose that $f(n) \ge c n^{\log_b a}$.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\geq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$= c n^{\log_b a} \log_b n$$

Hence,

$$T(n) = \mathbf{\Omega}(n^{\log_b a} \log_b n) \qquad \Rightarrow T(n) = \mathbf{\Omega}(n^{\log_b a} \log n).$$

60/530

Case 2. Now suppose that $f(n) \le c n^{\log_b a} (\log_b(n))^k$.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \cdot \left(\log_b \left(\frac{n}{b^i}\right)\right)^k$$

$$n = b^{\ell} \Rightarrow \ell = \log_b n$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell - 1} \left(\log_b \left(\frac{b^{\ell}}{b^i}\right)\right)^k$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell - 1} (\ell - i)^k$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell} i^k \sum_{i=1}^{\ell} i^{\ell + 1}$$

$$\approx \frac{c}{b} n^{\log_b a} \ell^{k+1} \qquad \Rightarrow T(n) = \mathcal{O}(n^{\log_b a} \log^{k+1} n).$$

Case 3. Now suppose that $f(n) \ge dn^{\log_b a + \epsilon}$, and that for sufficiently large n: $af(n/b) \le cf(n)$, for c < 1.

From this we get $a^i f(n/b^i) \le c^i f(n)$, where we assume that $n/b^{i-1} \ge n_0$ is still sufficiently large.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq \sum_{i=0}^{\log_b n - 1} c^i f(n) + \mathcal{O}(n^{\log_b a})$$

$$q < 1: \sum_{i=0}^n q^i = \frac{1 - q^{n+1}}{1 - q} \leq \frac{1}{1 - c} f(n) + \mathcal{O}(n^{\log_b a})$$

Hence,

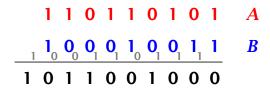
$$T(n) \le \mathcal{O}(f(n))$$

$$\Rightarrow T(n) = \Theta(f(n)).$$

Where did we use $f(n) \ge \Omega(n^{\log_b a + \epsilon})$?

Suppose we want to multiply two n-bit Integers, but our registers can only perform operations on integers of constant size.

For this we first need to be able to add two integers \mathbf{A} and \mathbf{B} :



This gives that two n-bit integers can be added in time $\mathcal{O}(n)$.

Suppose that we want to multiply an n-bit integer A and an m-bit integer B ($m \le n$).

	1	0	0	0	1	X	1	0	1	1	
						1	0	0	0	1	
 This is also nown as th method" for multiplying 					1	0	0	0	1	0	
 Note that the intermed bers that are generated 						0					
at most $m + n \le 2n$ bits			_1	0	0	0	1	0	0	0	
-			1	0	1	1	1	0	1	1	

Time requirement:

- ▶ Computing intermediate results: O(nm).
- ▶ Adding *m* numbers of length $\leq 2n$: $\mathcal{O}((m+n)m) = \mathcal{O}(nm)$.

A recursive approach:

Suppose that integers **A** and **B** are of length $n = 2^k$, for some k.



Then it holds that

$$A = A_1 \cdot 2^{\frac{n}{2}} + A_0$$
 and $B = B_1 \cdot 2^{\frac{n}{2}} + B_0$

Hence,

$$A \cdot B = A_1 B_1 \cdot 2^n + (A_1 B_0 + A_0 B_1) \cdot 2^{\frac{n}{2}} + A_0 B_0$$

Algorithm 3 $mult(A, B)$	
1: if $ A = B = 1$ then	$\mathcal{O}(1)$
2: return $a_0 \cdot b_0$	$\mathcal{O}(1)$
3: $splitA$ into A_0 and A_1	$\mathcal{O}(n)$
4: split B into B_0 and B_1	$\mathcal{O}(n)$
$5: Z_2 \leftarrow \operatorname{mult}(A_1, B_1)$	$T(\frac{n}{2})$
6: $Z_1 \leftarrow \text{mult}(A_1, B_0) + \text{mult}(A_0, B_1)$	$2T(\frac{n}{2}) + \mathcal{O}(n)$
7: $Z_0 \leftarrow \operatorname{mult}(A_0, B_0)$	$T(\frac{n}{2})$
8: return $Z_2 \cdot 2^n + Z_1 \cdot 2^{\frac{n}{2}} + Z_0$	$\mathcal{O}(n)$

We get the following recurrence:

$$T(n) = 4T\left(\frac{n}{2}\right) + \mathcal{O}(n) .$$

Master Theorem: Recurrence: $T[n] = aT(\frac{n}{b}) + f(n)$.

- ► Case 1: $f(n) = O(n^{\log_b a \epsilon})$ $T(n) = O(n^{\log_b a})$
- ► Case 2: $f(n) = \Theta(n^{\log_b a} \log^k n)$ $T(n) = \Theta(n^{\log_b a} \log^{k+1} n)$
- Case 3: $f(n) = \Omega(n^{\log_b a + \epsilon})$ $T(n) = \Theta(f(n))$

In our case a=4, b=2, and $f(n)=\Theta(n)$. Hence, we are in Case 1, since $n=\mathcal{O}(n^{2-\epsilon})=\mathcal{O}(n^{\log_b a-\epsilon})$.

We get a running time of $\mathcal{O}(n^2)$ for our algorithm.

⇒ Not better then the "school method".

We can use the following identity to compute Z_1 :

$$Z_1 = A_1 B_0 + A_0 B_1$$
 = Z_2 = Z_0
= $(A_0 + A_1) \cdot (B_0 + B_1) - A_1 B_1 - A_0 B_0$

Hence,

Algorithm 4 mult(A, B)1: **if** |A| = |B| = 1 then 2: **return** $a_0 \cdot b_0$ 3: split A into A_0 and A_1 4: split B into B_0 and B_1 6: $Z_0 \leftarrow \text{mult}(A_1, B_1)$ 7: $Z_1 \leftarrow \text{mult}(A_0 + A_1, B_0 + B_1) - Z_2 - Z_0$ 8: **return** $Z_2 \cdot 2^n + Z_1 \cdot 2^{\frac{n}{2}} + Z_0$ $T(\frac{n}{2})$ $T(\frac{n}{2})$ $T(\frac{n}{2}) + O(n)$

A more precise (correct) analysis would say that computing Z_1 needs time $T(\frac{n}{2}+1)+\mathcal{O}(n)$.

Example: Multiplying Two Integers

We get the following recurrence:

$$T(n) = 3T\left(\frac{n}{2}\right) + \mathcal{O}(n) .$$

Master Theorem: Recurrence: $T[n] = aT(\frac{n}{b}) + f(n)$.

- ► Case 1: $f(n) = \mathcal{O}(n^{\log_b a \epsilon})$ $T(n) = \Theta(n^{\log_b a})$
- ► Case 2: $f(n) = \Theta(n^{\log_b a} \log^k n)$ $T(n) = \Theta(n^{\log_b a} \log^{k+1} n)$
- Case 3: $f(n) = \Omega(n^{\log_b a + \epsilon})$ $T(n) = \Theta(f(n))$

Again we are in Case 1. We get a running time of $\Theta(n^{\log_2 3}) \approx \Theta(n^{1.59})$.

A huge improvement over the "school method".

6.3 The Characteristic Polynomial

Consider the recurrence relation:

$$c_0T(n) + c_1T(n-1) + c_2T(n-2) + \cdots + c_kT(n-k) = f(n)$$

This is the general form of a linear recurrence relation of order k with constant coefficients $(c_0, c_k \neq 0)$.

- ▶ T(n) only depends on the k preceding values. This means the recurrence relation is of order k.
- ▶ The recurrence is linear as there are no products of T[n]'s.
- If f(n) = 0 then the recurrence relation becomes a linear, homogenous recurrence relation of order k.

Note that we ignore boundary conditions for the moment.



6.3 The Characteristic Polynomial

Observations:

- ▶ The solution T[1], T[2], T[3], ... is completely determined by a set of boundary conditions that specify values for $T[1], \ldots, T[k]$.
- In fact, any k consecutive values completely determine the solution.
- k non-concecutive values might not be an appropriate set of boundary conditions (depends on the problem).

Approach:

- First determine all solutions that satisfy recurrence relation.
- Then pick the right one by analyzing boundary conditions.
- First consider the homogenous case.

The solution space

$$S = \left\{ \mathcal{T} = T[1], T[2], T[3], \dots \mid \mathcal{T} \text{ fulfills recurrence relation} \right\}$$

is a vector space. This means that if $\mathcal{T}_1, \mathcal{T}_2 \in S$, then also $\alpha \mathcal{T}_1 + \beta \mathcal{T}_2 \in S$, for arbitrary constants α, β .

How do we find a non-trivial solution?

We guess that the solution is of the form λ^n , $\lambda \neq 0$, and see what happens. In order for this guess to fulfill the recurrence we need

$$c_0\lambda^n + c_1\lambda^{n-1} + c_2 \cdot \lambda^{n-2} + \dots + c_k \cdot \lambda^{n-k} = 0$$

for all n > k.



Dividing by λ^{n-k} gives that all these constraints are identical to

$$\underbrace{c_0 \lambda^k + c_1 \lambda^{k-1} + c_2 \cdot \lambda^{k-2} + \dots + c_k}_{\text{characteristic polynomial } P[\lambda]} = 0$$

This means that if λ_i is a root (Nullstelle) of $P[\lambda]$ then $T[n] = \lambda_i^n$ is a solution to the recurrence relation.

Let $\lambda_1, \ldots, \lambda_k$ be the k (complex) roots of $P[\lambda]$. Then, because of the vector space property

$$\alpha_1\lambda_1^n + \alpha_2\lambda_2^n + \cdots + \alpha_k\lambda_k^n$$

is a solution for arbitrary values α_i .

Lemma 6

Assume that the characteristic polynomial has k distinct roots $\lambda_1, \ldots, \lambda_k$. Then all solutions to the recurrence relation are of the form

$$\alpha_1\lambda_1^n + \alpha_2\lambda_2^n + \cdots + \alpha_k\lambda_k^n$$
.

Proof.

There is one solution for every possible choice of boundary conditions for $T[1], \ldots, T[k]$.

We show that the above set of solutions contains one solution for every choice of boundary conditions.

Proof (cont.).

Suppose I am given boundary conditions T[i] and I want to see whether I can choose the $\alpha_i's$ such that these conditions are met:

Proof (cont.).

Suppose I am given boundary conditions T[i] and I want to see whether I can choose the $\alpha_i's$ such that these conditions are met:

$$\begin{pmatrix} \lambda_1 & \lambda_2 & \cdots & \lambda_k \\ \lambda_1^2 & \lambda_2^2 & \cdots & \lambda_k^2 \\ & \vdots & & \\ \lambda_1^k & \lambda_2^k & \cdots & \lambda_k^k \end{pmatrix} \begin{pmatrix} \alpha_1 \\ \alpha_2 \\ \vdots \\ \alpha_k \end{pmatrix} = \begin{pmatrix} T[1] \\ T[2] \\ \vdots \\ T[k] \end{pmatrix}$$

We show that the column vectors are linearly independent. Then the above equation has a solution.

$$\begin{vmatrix} \lambda_1 & \lambda_2 & \cdots & \lambda_{k-1} & \lambda_k \\ \lambda_1^2 & \lambda_2^2 & \cdots & \lambda_{k-1}^2 & \lambda_k^2 \\ \vdots & \vdots & & \vdots & \vdots \\ \lambda_1^k & \lambda_2^k & \cdots & \lambda_{k-1}^k & \lambda_k^k \end{vmatrix} = \prod_{i=1}^k \lambda_i \cdot \begin{vmatrix} 1 & 1 & \cdots & 1 & 1 \\ \lambda_1 & \lambda_2 & \cdots & \lambda_{k-1} & \lambda_k \\ \vdots & \vdots & & \vdots & \vdots \\ \lambda_1^{k-1} & \lambda_2^{k-1} & \cdots & \lambda_{k-1}^{k-1} & \lambda_k^{k-1} \end{vmatrix}$$

$$=\prod_{i=1}^k \lambda_i \cdot \begin{vmatrix} 1 & \lambda_1 & \cdots & \lambda_1^{k-2} & \lambda_1^{k-1} \\ 1 & \lambda_2 & \cdots & \lambda_2^{k-2} & \lambda_2^{k-1} \\ \vdots & \vdots & & \vdots & \vdots \\ 1 & \lambda_k & \cdots & \lambda_k^{k-2} & \lambda_k^{k-1} \end{vmatrix}$$

$$\begin{vmatrix} 1 & \lambda_{1} & \cdots & \lambda_{1}^{k-2} & \lambda_{1}^{k-1} \\ 1 & \lambda_{2} & \cdots & \lambda_{2}^{k-2} & \lambda_{2}^{k-1} \\ \vdots & \vdots & & \vdots & \vdots \\ 1 & \lambda_{k} & \cdots & \lambda_{k}^{k-2} & \lambda_{k}^{k-1} \end{vmatrix} = \\ \begin{vmatrix} 1 & \lambda_{1} - \lambda_{1} \cdot 1 & \cdots & \lambda_{1}^{k-2} - \lambda_{1} \cdot \lambda_{1}^{k-3} & \lambda_{1}^{k-1} - \lambda_{1} \cdot \lambda_{1}^{k-2} \\ 1 & \lambda_{2} - \lambda_{1} \cdot 1 & \cdots & \lambda_{2}^{k-2} - \lambda_{1} \cdot \lambda_{2}^{k-3} & \lambda_{2}^{k-1} - \lambda_{1} \cdot \lambda_{2}^{k-2} \\ \vdots & \vdots & & \vdots & \vdots \\ 1 & \lambda_{k} - \lambda_{1} \cdot 1 & \cdots & \lambda_{k}^{k-2} - \lambda_{1} \cdot \lambda_{k}^{k-3} & \lambda_{k}^{k-1} - \lambda_{1} \cdot \lambda_{k}^{k-2} \end{vmatrix}$$

$$\begin{vmatrix} 1 & \lambda_1 - \lambda_1 \cdot 1 & \cdots & \lambda_1^{k-2} - \lambda_1 \cdot \lambda_1^{k-3} & \lambda_1^{k-1} - \lambda_1 \cdot \lambda_1^{k-2} \\ 1 & \lambda_2 - \lambda_1 \cdot 1 & \cdots & \lambda_2^{k-2} - \lambda_1 \cdot \lambda_2^{k-3} & \lambda_2^{k-1} - \lambda_1 \cdot \lambda_2^{k-2} \\ \vdots & \vdots & & \vdots & & \vdots \\ 1 & \lambda_k - \lambda_1 \cdot 1 & \cdots & \lambda_k^{k-2} - \lambda_1 \cdot \lambda_k^{k-3} & \lambda_k^{k-1} - \lambda_1 \cdot \lambda_k^{k-2} \end{vmatrix} =$$

$$\begin{vmatrix} 1 & 0 & \cdots & 0 & 0 \\ 1 & (\lambda_2 - \lambda_1) \cdot 1 & \cdots & (\lambda_2 - \lambda_1) \cdot \lambda_2^{k-3} & (\lambda_2 - \lambda_1) \cdot \lambda_2^{k-2} \\ \vdots & \vdots & & \vdots & \vdots \\ 1 & (\lambda_k - \lambda_1) \cdot 1 & \cdots & (\lambda_k - \lambda_1) \cdot \lambda_k^{k-3} & (\lambda_k - \lambda_1) \cdot \lambda_k^{k-2} \end{vmatrix}$$

$$\begin{vmatrix} 1 & 0 & \cdots & 0 & 0 \\ 1 & (\lambda_2 - \lambda_1) \cdot 1 & \cdots & (\lambda_2 - \lambda_1) \cdot \lambda_2^{k-3} & (\lambda_2 - \lambda_1) \cdot \lambda_2^{k-2} \\ \vdots & \vdots & & \vdots & \vdots \\ 1 & (\lambda_k - \lambda_1) \cdot 1 & \cdots & (\lambda_k - \lambda_1) \cdot \lambda_k^{k-3} & (\lambda_k - \lambda_1) \cdot \lambda_k^{k-2} \end{vmatrix} =$$

$$\begin{vmatrix} \sum_{i=2}^{k} (\lambda_i - \lambda_1) \cdot \begin{pmatrix} 1 & \lambda_2 & \cdots & \lambda_2^{k-3} & \lambda_2^{k-2} \\ \vdots & \vdots & & \vdots & \vdots \\ 1 & \lambda_k & \cdots & \lambda_k^{k-3} & \lambda_k^{k-2} \end{pmatrix}$$

Repeating the above steps gives:

$$\begin{vmatrix} \lambda_1 & \lambda_2 & \cdots & \lambda_{k-1} & \lambda_k \\ \lambda_1^2 & \lambda_2^2 & \cdots & \lambda_{k-1}^2 & \lambda_k^2 \\ \vdots & \vdots & & \vdots & \vdots \\ \lambda_1^k & \lambda_2^k & \cdots & \lambda_{k-1}^k & \lambda_k^k \end{vmatrix} = \prod_{i=1}^k \lambda_i \cdot \prod_{i>\ell} (\lambda_i - \lambda_\ell)$$

Hence, if all λ_i 's are different, then the determinant is non-zero.

What happens if the roots are not all distinct?

Suppose we have a root λ_i with multiplicity (Vielfachheit) at least 2. Then not only is λ_i^n a solution to the recurrence but also $n\lambda_i^n$.

To see this consider the polynomial

$$P[\lambda] \cdot \lambda^{n-k} = c_0 \lambda^n + c_1 \lambda^{n-1} + c_2 \lambda^{n-2} + \dots + c_k \lambda^{n-k}$$

Since λ_i is a root we can write this as $Q[\lambda] \cdot (\lambda - \lambda_i)^2$. Calculating the derivative gives a polynomial that still has root λ_i .

This means

$$c_0 n \lambda_i^{n-1} + c_1 (n-1) \lambda_i^{n-2} + \dots + c_k (n-k) \lambda_i^{n-k-1} = 0$$

Hence,

$$c_0 \underbrace{n\lambda_i^n}_{T[n]} + c_1 \underbrace{(n-1)\lambda_i^{n-1}}_{T[n-1]} + \cdots + c_k \underbrace{(n-k)\lambda_i^{n-k}}_{T[n-k]} = 0$$

Suppose λ_i has multiplicity j. We know that

$$c_0 n \lambda_i^n + c_1 (n-1) \lambda_i^{n-1} + \dots + c_k (n-k) \lambda_i^{n-k} = 0$$

(after taking the derivative; multiplying with λ ; plugging in λ_i)

Doing this again gives

$$c_0 n^2 \lambda_i^n + c_1 (n-1)^2 \lambda_i^{n-1} + \dots + c_k (n-k)^2 \lambda_i^{n-k} = 0$$

We can continue j-1 times.

Hence, $n^{\ell} \lambda_i^n$ is a solution for $\ell \in 0, ..., j-1$.

Lemma 7

Let $P[\lambda]$ denote the characteristic polynomial to the recurrence

$$c_0T[n] + c_1T[n-1] + \cdots + c_kT[n-k] = 0$$

Let λ_i , $i=1,\ldots,m$ be the (complex) roots of $P[\lambda]$ with multiplicities ℓ_i . Then the general solution to the recurrence is given by

$$T[n] = \sum_{i=1}^{m} \sum_{j=0}^{\ell_i - 1} \alpha_{ij} \cdot (n^j \lambda_i^n) .$$

The full proof is omitted. We have only shown that any choice of α_{ij} 's is a solution to the recurrence.

Example: Fibonacci Sequence

$$T[0] = 0$$

 $T[1] = 1$
 $T[n] = T[n-1] + T[n-2]$ for $n \ge 2$

The characteristic polynomial is

$$\lambda^2 - \lambda - 1$$

Finding the roots, gives

$$\lambda_{1/2} = \frac{1}{2} \pm \sqrt{\frac{1}{4} + 1} = \frac{1}{2} \left(1 \pm \sqrt{5} \right)$$

Example: Fibonacci Sequence

Hence, the solution is of the form

$$\alpha \left(\frac{1+\sqrt{5}}{2}\right)^n + \beta \left(\frac{1-\sqrt{5}}{2}\right)^n$$

$$T[0] = 0$$
 gives $\alpha + \beta = 0$.

$$T[1] = 1$$
 gives

$$\alpha\left(\frac{1+\sqrt{5}}{2}\right)+\beta\left(\frac{1-\sqrt{5}}{2}\right)=1 \Longrightarrow \alpha-\beta=\frac{2}{\sqrt{5}}$$

Example: Fibonacci Sequence

Hence, the solution is

$$\frac{1}{\sqrt{5}} \left[\left(\frac{1+\sqrt{5}}{2} \right)^n - \left(\frac{1-\sqrt{5}}{2} \right)^n \right]$$

Consider the recurrence relation:

$$c_0T(n) + c_1T(n-1) + c_2T(n-2) + \cdots + c_kT(n-k) = f(n)$$

with $f(n) \neq 0$.

While we have a fairly general technique for solving homogeneous, linear recurrence relations the inhomogeneous case is different.

The general solution of the recurrence relation is

$$T(n) = T_h(n) + T_p(n) ,$$

where T_h is any solution to the homogeneous equation, and T_p is one particular solution to the inhomogeneous equation.

There is no general method to find a particular solution.

Example:

$$T[n] = T[n-1] + 1$$
 $T[0] = 1$

Then,

$$T[n-1] = T[n-2] + 1$$
 $(n \ge 2)$

Subtracting the first from the second equation gives,

$$T[n] - T[n-1] = T[n-1] - T[n-2]$$
 $(n \ge 2)$

or

$$T[n] = 2T[n-1] - T[n-2]$$
 $(n \ge 2)$

I get a completely determined recurrence if I add T[0] = 1 and T[1] = 2.

Example: Characteristic polynomial:

$$\underbrace{\lambda^2 - 2\lambda + 1}_{(\lambda - 1)^2} = 0$$

Then the solution is of the form

$$T[n] = \alpha 1^n + \beta n 1^n = \alpha + \beta n$$

$$T[0] = 1$$
 gives $\alpha = 1$.

$$T[1] = 2$$
 gives $1 + \beta = 2 \Rightarrow \beta = 1$.

If f(n) is a polynomial of degree r this method can be applied r+1 times to obtain a homogeneous equation:

$$T[n] = T[n-1] + n^2$$

Shift:

$$T[n-1] = T[n-2] + (n-1)^2 = T[n-2] + n^2 - 2n + 1$$

Difference:

$$T[n] - T[n-1] = T[n-1] - T[n-2] + 2n - 1$$

$$T[n] = 2T[n-1] - T[n-2] + 2n - 1$$

$$T[n] = 2T[n-1] - T[n-2] + 2n - 1$$

Shift:

$$T[n-1] = 2T[n-2] - T[n-3] + 2(n-1) - 1$$
$$= 2T[n-2] - T[n-3] + 2n - 3$$

Difference:

$$T[n] - T[n-1] = 2T[n-1] - T[n-2] + 2n - 1$$

- $2T[n-2] + T[n-3] - 2n + 3$

$$T[n] = 3T[n-1] - 3T[n-2] + T[n-3] + 2$$
 and so on...

Definition 8 (Generating Function)

Let $(a_n)_{n\geq 0}$ be a sequence. The corresponding

generating function (Erzeugendenfunktion) is

$$F(z) := \sum_{n \ge 0} a_n z^n ;$$

 exponential generating function (exponentielle Erzeugendenfunktion) is

$$F(z) := \sum_{n \ge 0} \frac{a_n}{n!} z^n .$$

Example 9

1. The generating function of the sequence (1, 0, 0, ...) is

$$F(z) = 1$$
.

2. The generating function of the sequence (1, 1, 1, ...) is

$$F(z) = \frac{1}{1-z}.$$

There are two different views:

A generating function is a formal power series (formale Potenzreihe).

Then the generating function is an algebraic object.

Let
$$f = \sum_{n \ge 0} a_n z^n$$
 and $g = \sum_{n \ge 0} b_n z^n$.

- **Equality:** f and g are equal if $a_n = b_n$ for all n.
- Addition: $f + g := \sum_{n \ge 0} (a_n + b_n) z^n$.
- ► Multiplication: $f \cdot g := \sum_{n\geq 0} c_n z^n$ with $c_n = \sum_{p=0}^n a_p b_{n-p}$.

There are no convergence issues here.

The arithmetic view:

We view a power series as a function $f: \mathbb{C} \to \mathbb{C}$.

Then, it is important to think about convergence/convergence radius etc.

What does $\sum_{n\geq 0} z^n = \frac{1}{1-z}$ mean in the algebraic view?

It means that the power series 1-z and the power series $\sum_{n\geq 0} z^n$ are invers, i.e.,

$$\left(1-z\right)\cdot\left(\sum_{n>0}^{\infty}z^n\right)=1.$$

This is well-defined.

Suppose we are given the generating function

$$\sum_{n\geq 0} z^n = \frac{1}{1-z} \cdot \begin{vmatrix} \frac{1}{(1-z)^2} \\ \text{Note that this requires a proof if we consider power series as algebraic} \end{vmatrix}$$

We can compute the derivative:

$$\sum_{n\geq 1} nz^{n-1} = \frac{1}{(1-z)^2}$$

$$\sum_{n\geq 0} (n+1)z^n$$

Hence, the generating function of the sequence $a_n = n + 1$ is $1/(1-z)^2$.

Formally the derivative of a formal !

power series $\sum_{n\geq 0} a_n z^n$ is defined as $\sum_{n\geq 0} na_n z^{n-1}$.

The known rules for differentiation

work for this definition. In particular, e.g. the derivative of $\frac{1}{1-7}$ is

consider power series as algebraic objects. However, we did not prove

this in the lecture.

We can repeat this

$$\sum_{n \ge 0} (n+1)z^n = \frac{1}{(1-z)^2} \ .$$

Derivative:

$$\underbrace{\sum_{n\geq 1} n(n+1)z^{n-1}}_{\sum_{n\geq 0} (n+1)(n+2)z^n} = \frac{2}{(1-z)^3}$$

Hence, the generating function of the sequence $a_n = (n+1)(n+2)$ is $\frac{2}{(1-z)^3}$.

Computing the k-th derivative of $\sum z^n$.

$$\sum_{n \ge k} n(n-1) \cdot \dots \cdot (n-k+1) z^{n-k} = \sum_{n \ge 0} (n+k) \cdot \dots \cdot (n+1) z^n$$
$$= \frac{k!}{(1-z)^{k+1}}.$$

Hence:

$$\sum_{n>0} \binom{n+k}{k} z^n = \frac{1}{(1-z)^{k+1}}.$$

The generating function of the sequence $a_n = \binom{n+k}{k}$ is $\frac{1}{(1-z)^{k+1}}$.

102/530

$$\sum_{n\geq 0} nz^n = \sum_{n\geq 0} (n+1)z^n - \sum_{n\geq 0} z^n$$
$$= \frac{1}{(1-z)^2} - \frac{1}{1-z}$$
$$= \frac{z}{(1-z)^2}$$

The generating function of the sequence $a_n = n$ is $\frac{z}{(1-z)^2}$.

We know

$$\sum_{n\geq 0} y^n = \frac{1}{1-y}$$

Hence,

$$\sum_{n\geq 0} a^n z^n = \frac{1}{1 - az}$$

The generating function of the sequence $f_n = a^n$ is $\frac{1}{1-az}$.

Suppose we have the recurrence $a_n = a_{n-1} + 1$ for $n \ge 1$ and $a_0 = 1$.

$$A(z) = \sum_{n\geq 0} a_n z^n$$

$$= a_0 + \sum_{n\geq 1} (a_{n-1} + 1) z^n$$

$$= 1 + z \sum_{n\geq 1} a_{n-1} z^{n-1} + \sum_{n\geq 1} z^n$$

$$= z \sum_{n\geq 0} a_n z^n + \sum_{n\geq 0} z^n$$

$$= zA(z) + \sum_{n\geq 0} z^n$$

$$= zA(z) + \frac{1}{1-z}$$

Solving for A(z) gives

$$\sum_{n\geq 0} a_n z^n = A(z) = \frac{1}{(1-z)^2} = \sum_{n\geq 0} (n+1) z^n$$

Hence, $a_n = n + 1$.

Some Generating Functions

n-th sequence element	generating function
1	$\frac{1}{1-z}$
n+1	$\frac{1}{(1-z)^2}$
$\binom{n+k}{k}$	$\frac{1}{(1-z)^{k+1}}$
n	$\frac{z}{(1-z)^2}$
a^n	$\frac{1}{1-az}$
n^2	$\frac{z(1+z)}{(1-z)^3}$
$\frac{1}{n!}$	e^z

Some Generating Functions

n-th sequence element	generating function
cf_n	cF
$f_n+{\cal g}_n$	F+G
$\sum_{i=0}^{n} f_i \mathcal{G}_{n-i}$	$F\cdot G$
f_{n-k} $(n \ge k)$; 0 otw.	$z^k F$
$\sum_{i=0}^{n} f_i$	$\frac{F(z)}{1-z}$
nf_n	$z\frac{\mathrm{d}F(z)}{\mathrm{d}z}$
$c^n f_n$	F(cz)

Solving Recursions with Generating Functions

- **1.** Set $A(z) = \sum_{n \ge 0} a_n z^n$.
- 2. Transform the right hand side so that boundary condition and recurrence relation can be plugged in.
- **3.** Do further transformations so that the infinite sums on the right hand side can be replaced by A(z).
- **4.** Solving for A(z) gives an equation of the form A(z) = f(z), where hopefully f(z) is a simple function.
- 5. Write f(z) as a formal power series. Techniques:
 - partial fraction decomposition (Partialbruchzerlegung)
 - lookup in tables
- **6.** The coefficients of the resulting power series are the a_n .



1. Set up generating function:

$$A(z) = \sum_{n \ge 0} a_n z^n$$

Transform right hand side so that recurrence can be plugged in:

$$A(z) = a_0 + \sum_{n>1} a_n z^n$$

2. Plug in:

$$A(z) = 1 + \sum_{n>1} (2a_{n-1})z^n$$

3. Transform right hand side so that infinite sums can be replaced by A(z) or by simple function.

$$A(z) = 1 + \sum_{n \ge 1} (2a_{n-1})z^n$$

$$= 1 + 2z \sum_{n \ge 1} a_{n-1}z^{n-1}$$

$$= 1 + 2z \sum_{n \ge 0} a_n z^n$$

$$= 1 + 2z \cdot A(z)$$

4. Solve for A(z).

$$A(z) = \frac{1}{1 - 2z}$$

5. Rewrite f(z) as a power series:

$$\sum_{n \geq 0} a_n z^n = A(z) = \frac{1}{1 - 2z} = \sum_{n \geq 0} 2^n z^n$$

1. Set up generating function:

$$A(z) = \sum_{n>0} a_n z^n$$

2./3. Transform right hand side:

$$A(z) = \sum_{n\geq 0} a_n z^n$$

$$= a_0 + \sum_{n\geq 1} a_n z^n$$

$$= 1 + \sum_{n\geq 1} (3a_{n-1} + n) z^n$$

$$= 1 + 3z \sum_{n\geq 1} a_{n-1} z^{n-1} + \sum_{n\geq 1} n z^n$$

$$= 1 + 3z \sum_{n\geq 0} a_n z^n + \sum_{n\geq 0} n z^n$$

$$= 1 + 3z A(z) + \frac{z}{(1-z)^2}$$

4. Solve for A(z):

$$A(z) = 1 + 3zA(z) + \frac{z}{(1-z)^2}$$

gives

$$A(z) = \frac{(1-z)^2 + z}{(1-3z)(1-z)^2} = \frac{z^2 - z + 1}{(1-3z)(1-z)^2}$$

5. Write f(z) as a formal power series:

We use partial fraction decomposition:

$$\frac{z^2 - z + 1}{(1 - 3z)(1 - z)^2} \stackrel{!}{=} \frac{A}{1 - 3z} + \frac{B}{1 - z} + \frac{C}{(1 - z)^2}$$

This gives

$$z^{2} - z + 1 = A(1 - z)^{2} + B(1 - 3z)(1 - z) + C(1 - 3z)$$

$$= A(1 - 2z + z^{2}) + B(1 - 4z + 3z^{2}) + C(1 - 3z)$$

$$= (A + 3B)z^{2} + (-2A - 4B - 3C)z + (A + B + C)$$

Example:
$$a_n = 3a_{n-1} + n$$
, $a_0 = 1$

5. Write f(z) as a formal power series:

This leads to the following conditions:

$$A + B + C = 1$$
$$2A + 4B + 3C = 1$$
$$A + 3B = 1$$

which gives

$$A = \frac{7}{4}$$
 $B = -\frac{1}{4}$ $C = -\frac{1}{2}$

5. Write f(z) as a formal power series:

$$A(z) = \frac{7}{4} \cdot \frac{1}{1 - 3z} - \frac{1}{4} \cdot \frac{1}{1 - z} - \frac{1}{2} \cdot \frac{1}{(1 - z)^2}$$

$$= \frac{7}{4} \cdot \sum_{n \ge 0} 3^n z^n - \frac{1}{4} \cdot \sum_{n \ge 0} z^n - \frac{1}{2} \cdot \sum_{n \ge 0} (n + 1) z^n$$

$$= \sum_{n \ge 0} \left(\frac{7}{4} \cdot 3^n - \frac{1}{4} - \frac{1}{2} (n + 1) \right) z^n$$

$$= \sum_{n \ge 0} \left(\frac{7}{4} \cdot 3^n - \frac{1}{2} n - \frac{3}{4} \right) z^n$$

6. This means $a_n = \frac{7}{4}3^n - \frac{1}{2}n - \frac{3}{4}$.

6.5 Transformation of the Recurrence

Example 10

$$f_0 = 1$$

$$f_1 = 2$$

$$f_n = f_{n-1} \cdot f_{n-2} \text{ for } n \ge 2 \ .$$

Define

$$g_n := \log f_n$$
.

Then

$$g_n = g_{n-1} + g_{n-2}$$
 for $n \ge 2$
 $g_1 = \log 2 = 1$ (for $\log = \log_2$), $g_0 = 0$
 $g_n = F_n$ (n -th Fibonacci number)
 $f_n = 2^{F_n}$

6.5 Transformation of the Recurrence

Example 11

$$f_1 = 1$$

 $f_n = 3f_{\frac{n}{2}} + n$; for $n = 2^k$, $k \ge 1$;

Define

$$g_k := f_{2^k}$$
.

Then:

$$g_0 = 1$$

 $g_k = 3g_{k-1} + 2^k, \ k \ge 1$

6 Recurrences

We get

$$g_k = 3 [g_{k-1}] + 2^k$$

$$= 3 [3g_{k-2} + 2^{k-1}] + 2^k$$

$$= 3^2 [g_{k-2}] + 32^{k-1} + 2^k$$

$$= 3^2 [3g_{k-3} + 2^{k-2}] + 32^{k-1} + 2^k$$

$$= 3^3 g_{k-3} + 3^2 2^{k-2} + 32^{k-1} + 2^k$$

$$= 2^k \cdot \sum_{i=0}^k \left(\frac{3}{2}\right)^i$$

$$= 2^k \cdot \frac{\left(\frac{3}{2}\right)^{k+1} - 1}{1/2} = 3^{k+1} - 2^{k+1}$$

6 Recurrences

Let
$$n = 2^k$$
:

$$g_k = 3^{k+1} - 2^{k+1}$$
, hence
 $f_n = 3 \cdot 3^k - 2 \cdot 2^k$
 $= 3(2^{\log 3})^k - 2 \cdot 2^k$
 $= 3(2^k)^{\log 3} - 2 \cdot 2^k$
 $= 3n^{\log 3} - 2n$.

6 Recurrences

Bibliography

[MS08] Kurt Mehlhorn, Peter Sanders:

Algorithms and Data Structures — The Basic Toolbox, Springer, 2008

Springer, 200

[CLRS90] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein:

Introduction to algorithms (3rd ed.), MIT Press and McGraw-Hill, 2009

[Liu85] Chung Laung Liu:

Elements of Discrete Mathematics

McGraw-Hill, 1985

The Karatsuba method can be found in [MS08] Chapter 1. Chapter 4.3 of [CLRS90] covers the "Substitution method" which roughly corresponds to "Guessing-induction". Chapters 4.4, 4.5, 4.6 of this book cover the master theorem. Methods using the characteristic polynomial and generating functions can be found in [Liu85] Chapter 10.

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.

Dynamic Set Operations

- ▶ *S*. search(*k*): Returns pointer to object x from S with key[x] = k or null.
- S. insert(x): Inserts object x into set S. key[x] must not currently exist in the data-structure.
- S. delete(x): Given pointer to object x from S, delete x from the set.
- S. minimum(): Return pointer to object with smallest key-value in S.
- S. maximum(): Return pointer to object with largest key-value in S.
- S. successor(x): Return pointer to the next larger element in S or null if x is maximum.
- S. predecessor(x): Return pointer to the next smaller element in S or null if x is minimum.



Dynamic Set Operations

- ▶ S. union(S'): Sets $S := S \cup S'$. The set S' is destroyed.
- ▶ S. merge(S'): Sets $S := S \cup S'$. Requires $S \cap S' = \emptyset$.
- ► *S.* split(k, S'): $S := \{x \in S \mid \text{key}[x] \le k\}, S' := \{x \in S \mid \text{key}[x] > k\}.$
- ► *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*.
- ▶ *S.* empty(): Tell if *S* contains any object.

Queue:

- S. enqueue(x): Insert an element.
- S. dequeue(): Return the element that is longest in the structure; delete it from S.
- ▶ *S.* empty(): Tell if *S* contains any object.

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:

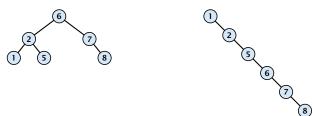
- S. insert(x): Insert an element x.
- S. delete(x): Delete the element pointed to by x.
- ▶ *S.* search(k): Return a pointer to an element e with key[e] = k in S if it exists; otherwise return null.

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:

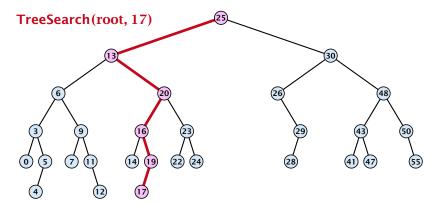


7.1 Binary Search Trees

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

- ightharpoonup T. insert(x)
- ightharpoonup T. delete(x)
- ightharpoonup T. search(k)
- ightharpoonup T. successor(x)
- ightharpoonup T. predecessor(x)
- ightharpoonup T. minimum()
- ightharpoonup T. maximum()

Binary Search Trees: Searching



Algorithm 1 TreeSearch(x, k)

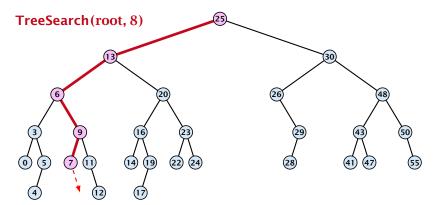
1: **if** x = null or k = key[x] **return** x

2: **if** k < key[x] **return** TreeSearch(left[x], k)

3: **else return** TreeSearch(right[x], k)



Binary Search Trees: Searching



Algorithm 1 TreeSearch(x, k)

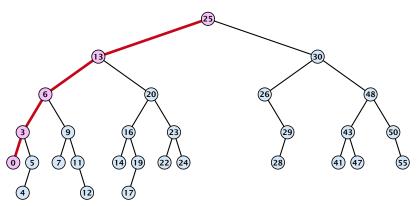
1: **if** x = null or k = key[x] **return** x

2: **if** k < key[x] **return** TreeSearch(left[x], k)

3: **else return** TreeSearch(right[x], k)



Binary Search Trees: Minimum

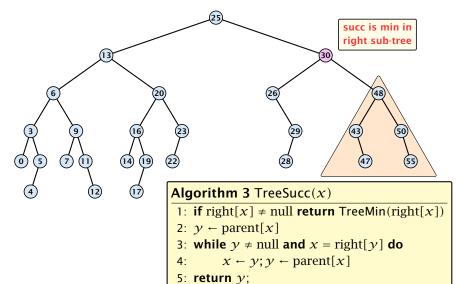


Algorithm 2 TreeMin(x)

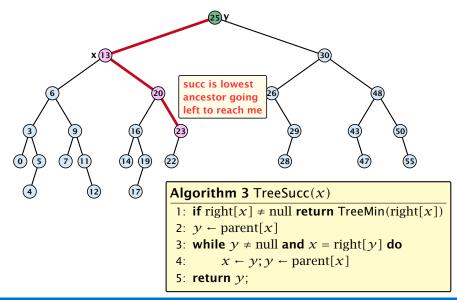
- 1: **if** x = null or left[x] = null return x
- 2: **return** TreeMin(left[x])



Binary Search Trees: Successor

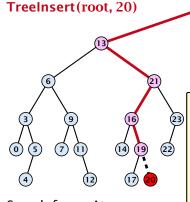


Binary Search Trees: Successor



Binary Search Trees: Insert

Insert element not in the tree.

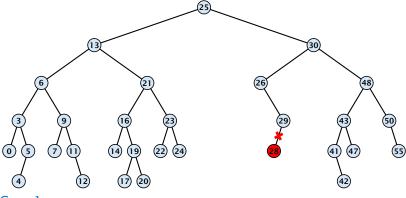


Search for z. At some point the search stops at a null-pointer. This is the place to insert z.

```
Algorithm 4 TreeInsert(x,z)
```

- 1: if x = null then
 - 2: $root[T] \leftarrow z$; parent[z] \leftarrow null;
- 3: return,
- 4: **if** key[x] > key[z] **then**
- 5: **if** left[x] = null **then**
- 6: $\operatorname{left}[x] \leftarrow z$; parent $[z] \leftarrow x$;
- 7: **else** Treelnsert(left[x], z); 8: **else**
- 9: **if** right[x] = null **then**
- 10: $\operatorname{right}[x] \leftarrow z$; parent $[z] \leftarrow x$;
- 11: **else** TreeInsert(right[x], z);

Binary Search Trees: Delete

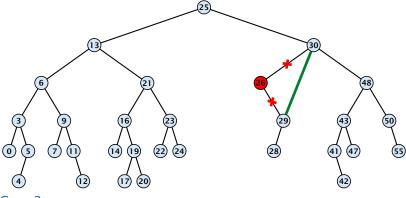


Case 1:

Element does not have any children

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

Binary Search Trees: Delete

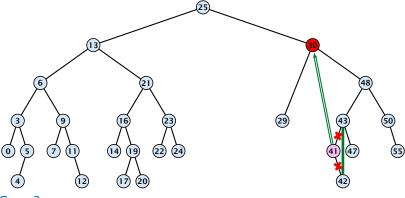


Case 2:

Element has exactly one child

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

Binary Search Trees: Delete



Case 3:

Element has two children

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

Binary Search Trees: Delete

```
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
```

137/530

Balanced Binary Search Trees

All operations on a binary search tree can be performed in time $\mathcal{O}(h)$, where h denotes the height of the tree.

However the height of the tree may become as large as $\Theta(n)$.

Balanced Binary Search Trees

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.



Binary Search Trees (BSTs)

Bibliography

[MS08] Kurt Mehlhorn, Peter Sanders:

Algorithms and Data Structures — The Basic Toolbox,

Springer, 2008

[CLRS90] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein:

Introduction to Algorithms (3rd ed.), MIT Press and McGraw-Hill, 2009

Binary search trees can be found in every standard text book. For example Chapter 7.1 in [MS08] and Chapter 12 in [CLRS90].

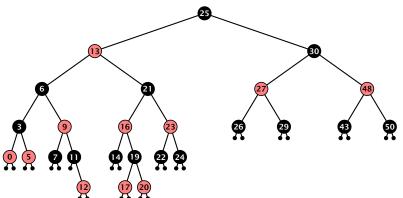
Definition 12

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.
- 2. All leaf nodes are black.
- 3. For each node, all paths to descendant leaves contain the same number of black nodes.
- 4. If a node is red then both its children are black.

The null-pointers in a binary search tree are replaced by pointers to special null-vertices, that do not carry any object-data

Red Black Trees: Example



Lemma 13

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

Definition 14

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 15

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 15.

Induction on the height of v.

base case (height(v) = 0)

- ▶ If height(v) (maximum distance btw. v and a node in the sub-tree rooted at v) is 0 then v is a leaf.
- ▶ The black height of v is 0.
- ► The sub-tree rooted at v contains $0 = 2^{bh(v)} 1$ inner vertices.

Proof (cont.)

induction step

- Supose v is a node with height(v) > 0.
- $\triangleright v$ has two children with strictly smaller height.
- ▶ These children (c_1, c_2) either have $bh(c_i) = bh(v)$ or $bh(c_{i}) = bh(v) - 1.$
- By induction hypothesis both sub-trees contain at least $2^{\mathrm{bh}(v)-1}-1$ internal vertices.
- ► Then T_v contains at least $2(2^{bh(v)-1}-1)+1 \ge 2^{bh(v)}-1$ vertices.



143/530

Proof of Lemma 13.

Let h denote the height of the red-black tree, and let P denote a path from the root to the furthest leaf.

At least half of the node on P must be black, since a red node must be followed by a black node.

Hence, the black height of the root is at least h/2.

The tree contains at least $2^{h/2}-1$ internal vertices. Hence, $2^{h/2}-1 \le n$.

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



Definition 1

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

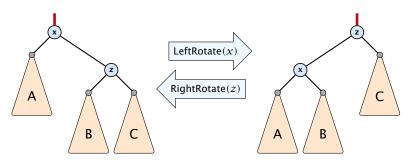
- 1. The root is black.
- 2. All leaf nodes are black.
- 3. For each node, all paths to descendant leaves contain the same number of black nodes.
- 4. If a node is red then both its children are black.

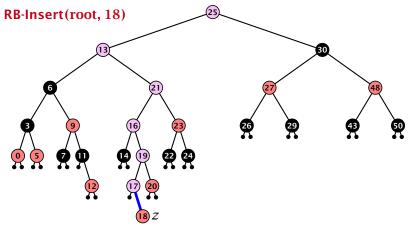
The null-pointers in a binary search tree are replaced by pointers to special null-vertices, that do not carry any object-data.

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:





Insert:

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



Invariant of the fix-up algorithm:

- z is a red node
- the black-height property is fulfilled at every node
- the only violation of red-black properties occurs at z and parent[z]
 - 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.

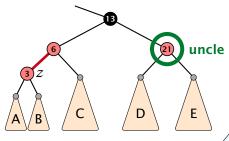
12. Jan. 2025

```
Algorithm 10 InsertFix(z)
 1: while parent[z] \neq null and col[parent[z]] = red do
         if parent[z] = left[gp[z]] then z in left subtree of grandparent
 2:
 3:
              uncle \leftarrow right[grandparent[z]]
              if col[uncle] = red then
 4:
                                                             Case 1: uncle red
                  col[p[z]] \leftarrow black; col[u] \leftarrow black;
 5:
                  col[gp[z]] \leftarrow red; z \leftarrow grandparent[z];
 6:
 7:
             else
                                                           Case 2: uncle black
                  if z = right[parent[z]] then
 8:
                                                              2a: z right child
                       z \leftarrow p[z]; LeftRotate(z);
 9:
                  col[p[z]] \leftarrow black; col[gp[z]] \leftarrow red; 2b: z left child
10:
11:
                  RightRotate(gp[z]);
12:
         else same as then-clause but right and left exchanged
13: col(root[T]) \leftarrow black;
```

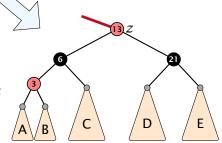


12. Jan. 2025

Case 1: Red Uncle

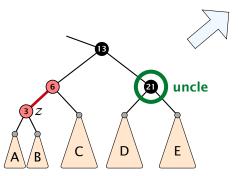


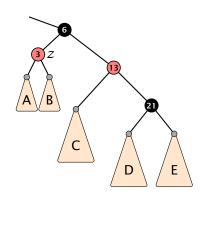
- 1. recolour
- 2. move z to grand-parent
- 3. invariant is fulfilled for new z
- 4. you made progress



Case 2b: Black uncle and z is left child

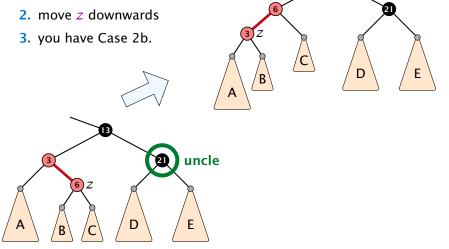
- 1. rotate around grandparent
- re-colour to ensure that black height property holds
- 3. you have a red black tree





Case 2a: Black uncle and z is right child

1. rotate around parent



Running time:

- ▶ Only Case 1 may repeat; but only h/2 many steps, where h is the height of the tree.
- Case 2a → Case 2b → red-black tree
- Case 2b → red-black tree

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.

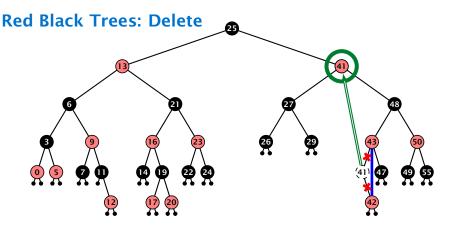
Red Black Trees: Delete

First do a standard delete.

If the spliced out node x was red everything is fine.

If it was black there may be the following problems.

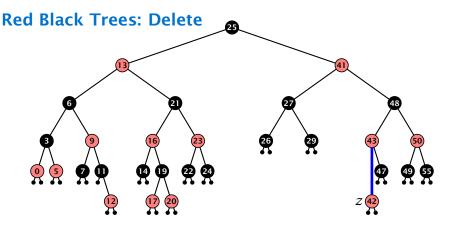
- Parent and child of x were red; two adjacent red vertices.
- If you delete the root, the root may now be red.
- Every path from an ancestor of x to a descendant leaf of x changes the number of black nodes. Black height property might be violated.



Case 3:

Element has two children

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



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.

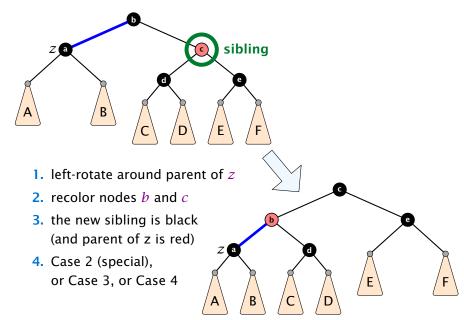
Red Black Trees: Delete

Invariant of the fix-up algorithm

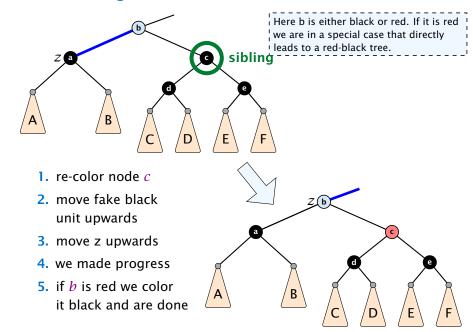
- ► the node z is black
- if we "assign" a fake black unit to the edge from z to its parent then the black-height property is fulfilled

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

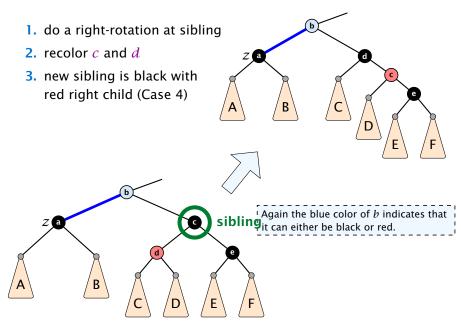
Case 1: Sibling of z is red



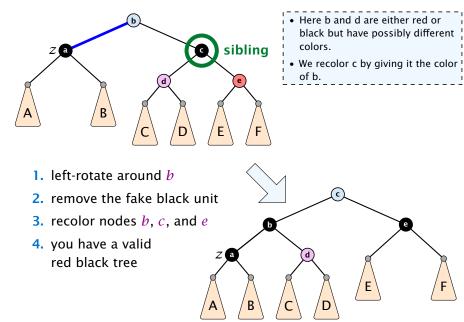
Case 2: Sibling is black with two black children



Case 3: Sibling black with one black child to the right



Case 4: Sibling is black with red right child



Running time:

- only Case 2 can repeat; but only h many steps, where h is the height of the tree
- Case 1 → Case 2 (special) → red black tree Case 1 → Case 3 → Case 4 → red black tree Case 1 → Case 4 → red black tree
- Case 3 → Case 4 → red black tree
- Case 4 → red black tree

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.

Red-Black Trees

Bibliography

[CLRS90] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein: Introduction to Algorithms (3rd ed.), MIT Press and McGraw-Hill, 2009

Red black trees are covered in detail in Chapter 13 of [CLRS90].

Disadvantage of balanced search trees:

- worst case; no advantage for easy inputs
- additional memory required
- complicated implementation

Splay Trees:

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

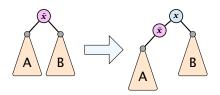


find(x)

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

insert(x)

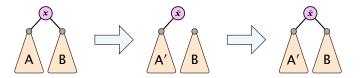
- search for x; \bar{x} is last visited element during search (successer or predecessor of x)
- splay(\bar{x}) moves \bar{x} to the root
- insert x as new root



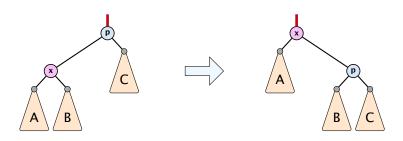
The illustration shows the case when \bar{x} is the predecessor of x.

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

Splay: Zig Case

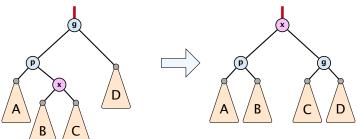


better option splay(x):

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

Note that moveToRoot(x) does the same.

Splay: Zigzag Case

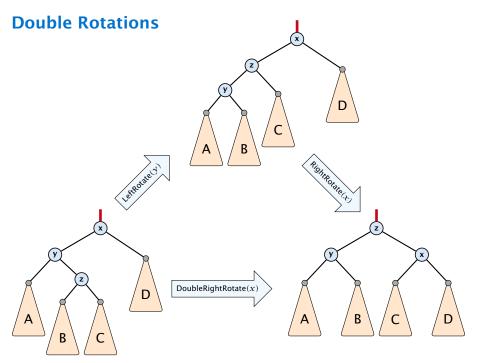


better option splay(x):

- 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)

Note that moveToRoot(x) does the same.



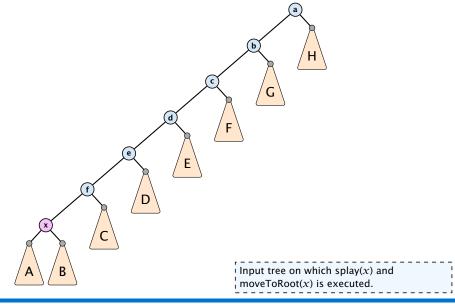


Splay: Zigzig Case This case is different between moveToRoot(x) and splay(x).

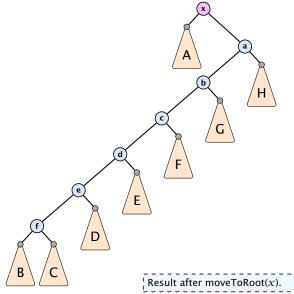
better option splay(x):

- zigzig case: if x is left child and parent of x is left child (or x right child, parent of x right child)
- do right roation around grand-parent followed by right rotation around parent (resp. left rotations)

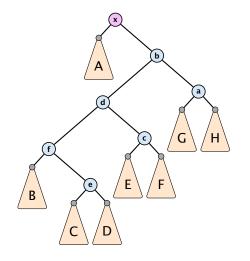
Splay vs. Move to Root



Splay vs. Move to Root



Splay vs. Move to Root



Result after splay(x).

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 $\mathcal{O}(\cos t(T_{\min}))$, where T_{\min} is an optimal static search tree.

 $\operatorname{depth}_T(x)$ is the number of edges on a path from the root of T to x.

Theorem given without proof.



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 16

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

Amortized Analysis

Definition 17

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 $\sum_i k_i \cdot t_i(n)$.

12. Jan. 2025

Potential Method

Introduce a potential for the data structure.

- $\Phi(D_i)$ is the potential after the *i*-th operation.
- ▶ Amortized cost of the i-th operation is

$$\hat{c}_i = c_i + \Phi(D_i) - \Phi(D_{i-1}) \ . \label{eq:constraint}$$

▶ Show that $\Phi(D_i) \ge \Phi(D_0)$.

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.

Example: Stack

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.

Actual cost:

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

Example: Stack

Use potential function $\Phi(S)$ = number of elements on the stack.

Amortized cost:

► S. push(): cost

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

▶ *S.* pop(): cost

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

Note that the analysis becomes wrong if pop() or multipop() are called on an empty stack.

 \triangleright S. multipop(k): cost

$$\hat{C}_{mn} = C_{mn} + \Delta \Phi = \min\{\text{size}, k\} - \min\{\text{size}, k\} \le 0$$
.

Example: Binary Counter

Incrementing a binary counter:

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

Incrementing an n-bit binary counter may require to examine n-bits, and maybe change them.

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).

Example: Binary Counter

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

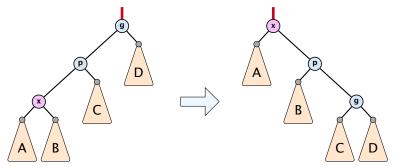


$$\Delta\Phi = r'(x) + r'(p) - r(x) - r(p)$$
$$= r'(p) - r(x)$$
$$\leq r'(x) - r(x)$$

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

Splay: Zigzig Case

Last inequality follows from next slide.



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

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

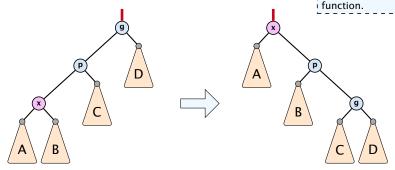
$$\leq r'(x) + r'(g) - r(x) - r(x)$$

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

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

$$\leq -2 + 3(r'(x) - r(x)) \Rightarrow \text{cost}_{\text{zigzig}} \leq 3(r'(x) - r(x))$$

Splay: Zigzig Case



The last inequality holds

because log is a concave

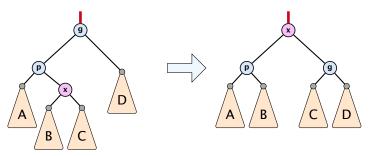
$$\frac{1}{2} \left(r(x) + r'(g) - 2r'(x) \right)$$

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

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

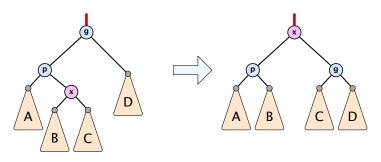
$$\leq \log \left(\frac{1}{2} \frac{s(x)}{s'(x)} + \frac{1}{2} \frac{s'(g)}{s'(x)} \right) \leq \log \left(\frac{1}{2} \right) = -1$$

Splay: Zigzag Case



$$\begin{split} \Delta \Phi &= r'(x) + r'(p) + r'(g) - r(x) - r(p) - r(g) \\ &= r'(p) + r'(g) - r(x) - r(p) \\ &\leq r'(p) + r'(g) - r(x) - r(x) \\ &= r'(p) + r'(g) - 2r'(x) + 2r'(x) - 2r(x) \\ &\leq -2 + 2(r'(x) - r(x)) \quad \Rightarrow \operatorname{cost}_{\mathsf{zigzag}} \leq 3(r'(x) - r(x)) \end{split}$$

Splay: Zigzag Case



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

Amortized cost of the whole splay operation:

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

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

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

The first one is added due to the fact that so far for each step of a splay-operation we have only counted the number of rotations, but the cost is 1+#rotations.

The second one comes from the zig-operation. Note that we have at most one zig-operation during a splay.



Splay Trees

Bibliography

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.

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

How to augment a data-structure

- choose an underlying data-structure
- determine additional information to be stored in the underlying structure
- 3. verify/show how the additional information can be maintained for the basic modifying operations on the underlying structure.
- 4. develop the new operations
- 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.

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

- 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 $\mathcal{O}(\log n)$.

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

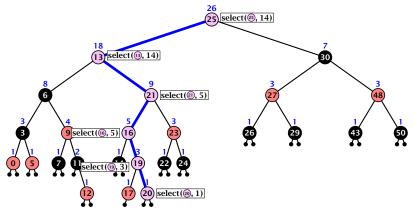
5: **return** Select(left[x], i)

6: **else**

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



Select(x, i)



Find-by-rank:

- 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



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

3. How do we maintain information?

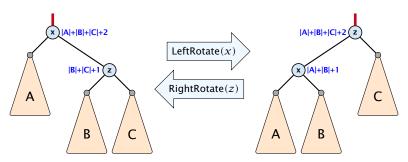
Search(k): Nothing to do.

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.

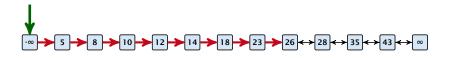
Bibliography

[CLRS90] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein: Introduction to Algorithms (3rd ed.), MIT Press and McGraw-Hill, 2009

See Chapter 14 of [CLRS90].

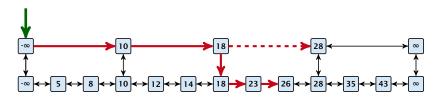
Why do we not use a list for implementing the ADT Dynamic Set?

- \blacktriangleright time for search $\Theta(n)$
- time for insert $\Theta(n)$ (dominated by searching the item)
- time for delete $\Theta(1)$ if we are given a handle to the object, otw. $\Theta(n)$



How can we improve the search-operation?

Add an express lane:



Let $|L_1|$ denote the number of elements in the "express lane", and $|L_0|=n$ the number of all elements (ignoring dummy elements).

Worst case search time: $|L_1| + \frac{|L_0|}{|L_1|}$ (ignoring additive constants)

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

Add more express lanes. Lane L_i contains roughly every $\frac{L_{i-1}}{L_i}$ -th item from list L_{i-1} .

Search(x) $(k + 1 \text{ lists } L_0, \ldots, L_k)$

- Find the largest item in list L_k that is smaller than x. At most $|L_k| + 2$ steps.
- Find the largest item in list L_{k-1} that is smaller than x. At most $\lceil \frac{|L_{k-1}|}{|L_k|+1} \rceil + 2$ steps.
- Find the largest item in list L_{k-2} that is smaller than x. At most $\left\lceil \frac{|L_{k-2}|}{|L_{k-1}|+1} \right\rceil + 2$ steps.
- **.**..
- ► At most $|L_k| + \sum_{i=1}^k \frac{L_{i-1}}{L_i} + 3(k+1)$ steps.

12. Jan. 2025

Choose ratios between list-lengths evenly, i.e., $\frac{|L_{i-1}|}{|L_i|} = r$, and, hence, $L_k \approx r^{-k}n$.

Worst case running time is: $\mathcal{O}(r^{-k}n + kr)$. Choose $r = n^{\frac{1}{k+1}}$. Then

$$r^{-k}n + kr = \left(n^{\frac{1}{k+1}}\right)^{-k}n + kn^{\frac{1}{k+1}}$$
$$= n^{1-\frac{k}{k+1}} + kn^{\frac{1}{k+1}}$$
$$= (k+1)n^{\frac{1}{k+1}}.$$

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

How to do insert and delete?

If we want that in L_i we always skip over roughly the same number of elements in L_{i-1} an insert or delete may require a lot of re-organisation.

Use randomization instead!

Insert:

- A search operation gives you the insert position for element x in every list.
- ▶ Flip a coin until it shows head, and record the number $t \in \{1, 2, ...\}$ of trials needed.
- ▶ Insert x into lists L_0, \ldots, L_{t-1} .

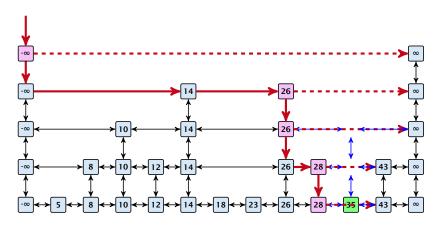
Delete:

- You get all predecessors via backward pointers.
- Delete x in all lists it actually appears in.

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



Insert (35):





High Probability

Definition 18 (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 $1 - \frac{1}{n^{\alpha}}$.

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



High Probability

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

Then the probability that all E_i hold is at least

$$\Pr[E_1 \wedge \cdots \wedge E_{\ell}] = 1 - \Pr[\bar{E}_1 \vee \cdots \vee \bar{E}_{\ell}]$$

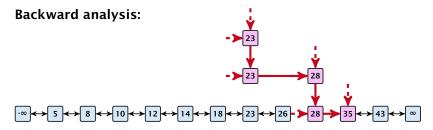
$$\geq 1 - n^c \cdot n^{-\alpha}$$

$$= 1 - n^{c - \alpha}.$$

This means $E_1 \wedge \cdots \wedge E_\ell$ holds with high probability.

Lemma 19

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.).



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.



7.5 Skip Lists 12. Jan. 2025

Estimation for Binomial Coefficients

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

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

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

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 .

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.

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

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

$$\leq \left(\frac{2e(\beta + \alpha)}{2^\beta}\right)^k n^{-\alpha}$$

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 $z = 7\alpha y \log n$, $\alpha \ge 1$.

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.

Skip Lists

Bibliography

[GT98] Michael T. Goodrich, Roberto Tamassia Data Structures and Algorithms in JAVA, John Wiley, 1998

Skip lists are covered in Chapter 7.5 of [GT98].

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:

- 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.

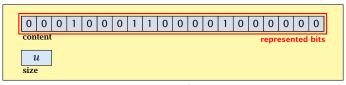


212/530

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 5 array.min()

```
1: for (i = 0; i < \text{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)

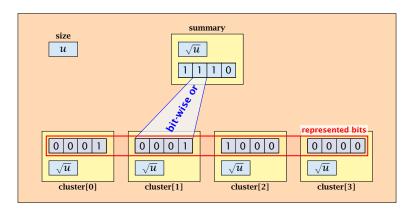
1: for (i = x + 1; i < size; i++) do 2: if content[i] = 1 then return i; 3: return null;

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 $\left\lfloor \frac{x}{\sqrt{u}} \right\rfloor$.

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 9 insert(x)

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

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: $offs \leftarrow cluster[maxcluster]. max()$
- 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 \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();

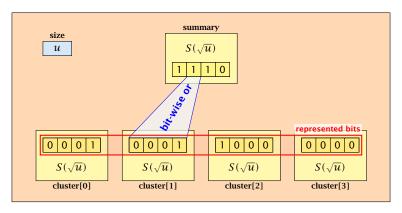
6: return predcluster ∘ offs;

7: return null;
```

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

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.

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: cluster[high(x)].insert(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;
```

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

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

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 $T_{\text{mem}}(u) = \mathcal{O}(\log \log u)$.

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

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

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

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

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 $T_{\text{min}}(u)$.

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

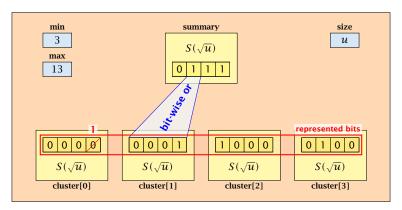
Set $\ell := \log u$ and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

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

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

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).



Advantages of having max/min pointers:

- 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:
T_{\text{succ}}(u) = T_{\text{succ}}(\sqrt{u}) + 1 \Rightarrow T_{\text{succ}}(u) = \mathcal{O}(\log \log u).
```

240/530

```
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
       if x = \min then
                                            find new minimum
            firstcluster \leftarrow summary.min();
            offs ← cluster[firstcluster]. min();
6:
      x \leftarrow firstcluster \circ offs;
 7:
         \min \leftarrow x;
        cluster[high(x)]. delete(low(x));
 9:
                                                      delete
                         continued...
```

```
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 $T_{\text{del}}(u) = \mathcal{O}(\log \log u)$.

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 $R(k^2) \le k^2 2$ for $k^2 \ge 4$.
 - Obviously, this holds for $k^2 = 4$.
 - For $k^2 > 4$ we have

$$R(k^{2}) = (1+k)R(k) + ck$$

$$\leq (1+k)(k-2) + k \leq k^{2} - 2$$

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

van Emde Boas Trees

Bibliography

[CLRS90] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein: Introduction to Algorithms (3rd ed.), MIT Press and McGraw-Hill, 2009

See Chapter 20 of [CLRS90].

7.7 Hashing

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.

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.

7.7 Hashing

Definitions:

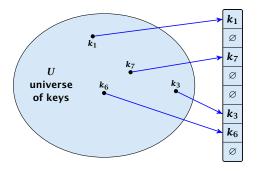
- ▶ Universe U of keys, e.g., $U \subseteq \mathbb{N}_0$. U very large.
- ▶ Set $S \subseteq U$ of keys, $|S| = m \le |U|$.
- Array $T[0, \ldots, n-1]$ hash-table.
- ► Hash function $h: U \rightarrow [0, ..., n-1]$.

The hash-function h should fulfill:

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

Direct Addressing

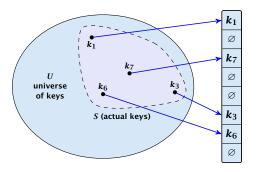
Ideally the hash function maps all keys to different memory locations.



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.

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.



12. Jan. 2025

Typically, collisions do not appear once the size of the set S of actual keys gets close to n, but already when $|S| \ge \omega(\sqrt{n})$.

Lemma 20

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

$$1 - e^{-\frac{m(m-1)}{2n}} \approx 1 - e^{-\frac{m^2}{2n}} .$$

Uniform hashing:

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



12. Jan. 2025

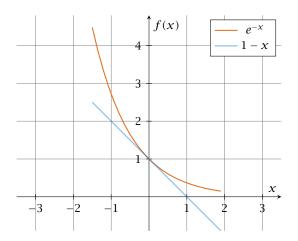
Proof.

Let $A_{m,n}$ denote the event that inserting m keys into a table of size n does not generate a collision. Then

$$\Pr[A_{m,n}] = \prod_{\ell=1}^{m} \frac{n-\ell+1}{n} = \prod_{j=0}^{m-1} \left(1 - \frac{j}{n}\right)$$

$$\leq \prod_{j=0}^{m-1} e^{-j/n} = e^{-\sum_{j=0}^{m-1} \frac{j}{n}} = e^{-\frac{m(m-1)}{2n}}.$$

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.



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



Resolving Collisions

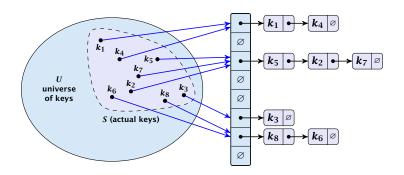
The methods for dealing with collisions can be classified into the two main types

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

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.



12. Jan. 2025 257/530

Let A denote a strategy for resolving collisions. We use the following notation:

- A⁺ denotes the average time for a successful search when using A;
- ▶ A^- denotes the average time for an **unsuccessful** search when using A;
- We parameterize the complexity results in terms of $\alpha := \frac{m}{n}$, the so-called fill factor of the hash-table.

We assume uniform hashing for the following analysis.

The time required for an unsuccessful search is 1 plus the length of the list that is examined. The average length of a list is $\alpha = \frac{m}{n}$. Hence, if A is the collision resolving strategy "Hashing with Chaining" we have

$$A^- = 1 + \alpha .$$

7.7 Hashing 12. Jan. 2025 259/530

For a successful search observe that we do **not** choose a list at random, but we consider a random key k in the hash-table and ask for the search-time for k.

This is 1 plus the number of elements that lie before k in k's list.

Let k_{ℓ} denote the ℓ -th key inserted into the table.

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, $\Pr[X_{ij}=1]=1/n$ for uniform hashing.

The expected successful search cost is

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



260/530

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

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

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.



Open Addressing

All objects are stored in the table itself.

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 $0,\ldots,n-1$.

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.

Open Addressing

Choices for h(k, j):

Linear probing:

$$h(k,i) = h(k) + i \mod n$$

(sometimes: $h(k,i) = h(k) + ci \mod n$).

- 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).

12. Jan. 2025

264/530

Linear Probing

- Advantage: Cache-efficiency. The new probe position is very likely to be in the cache.
- Disadvantage: Primary clustering. Long sequences of occupied table-positions get longer as they have a larger probability to be hit. Furthermore, they can merge forming larger sequences.

Lemma 21

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.
- Secondary clustering: caused by the fact that all keys mapped to the same position have the same probe sequence.

Lemma 22

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.

Lemma 23

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

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

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

Open Addressing

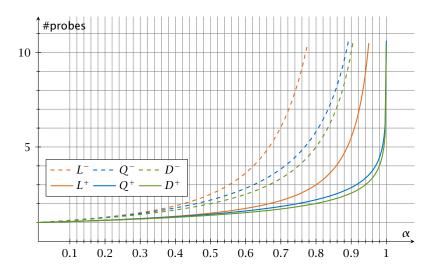
Some values:

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



12. Jan. 2025

Open Addressing





12. Jan. 2025 269/530

Analysis of Idealized Open Address Hashing

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).

Analysis of Idealized Open Address Hashing

Let X denote a random variable describing the number of probes in an unsuccessful search.

Let A_i denote the event that the i-th probe occurs and is to a non-empty slot.

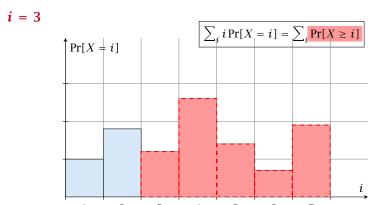
$$Pr[A_1 \cap A_2 \cap \cdots \cap A_{i-1}]$$

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

$$\Pr[X \ge i] = \frac{m}{n} \cdot \frac{m-1}{n-1} \cdot \frac{m-2}{n-2} \cdot \dots \cdot \frac{m-i+2}{n-i+2}$$
$$\le \left(\frac{m}{n}\right)^{i-1} = \alpha^{i-1} .$$

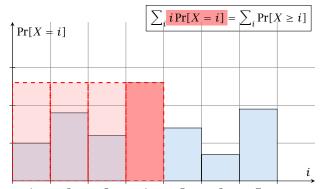
$$E[X] = \sum_{i=1}^{\infty} \Pr[X \ge i] \le \sum_{i=1}^{\infty} \alpha^{i-1} = \sum_{i=0}^{\infty} \alpha^{i} = \frac{1}{1-\alpha}.$$

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



The j-th rectangle appears in both sums j^6 times. (j times in the first due to multiplication with j; and j times in the second for summands $i = 1, 2, \dots, j$





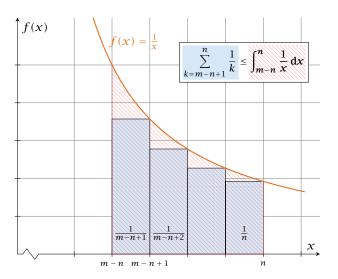
The j-th rectangle² appears in both sums j⁶ times. (j times in the first due to multiplication with j; and j times in the second for summands i = 1, 2, ..., j)

The number of probes in a successful search for k is equal to the number of probes made in an unsuccessful search for k at the time that k is inserted.

Let k be the i+1-st element. The expected time for a search for k is at most $\frac{1}{1-i/n}=\frac{n}{n-i}$.

$$\frac{1}{m} \sum_{i=0}^{m-1} \frac{n}{n-i} = \frac{n}{m} \sum_{i=0}^{m-1} \frac{1}{n-i} = \frac{1}{\alpha} \sum_{k=n-m+1}^{n} \frac{1}{k}$$

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



Deletions in Hashtables

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.

Deletions in Hashtables

- Simply removing a key might interrupt the probe sequence of other keys which then cannot be found anymore.
- One can delete an element by replacing it with a deleted-marker.
 - 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.



12. Jan. 2025 277/530

Deletions for Linear Probing

- 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.

Deletions for Linear Probing

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: $p \leftarrow \text{succ}(p)$ 7: insert(y)

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

Pointers into the hash-table become invalid.



Regardless, of the choice of hash-function there is always an input (a set of keys) that has a very poor worst-case behaviour.

Therefore, so far we assumed that the hash-function is random so that regardless of the input the average case behaviour is good.

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 $\mathcal H$.



Definition 24

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} .

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

Definition 25

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 :

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

This requirement clearly implies a universal hash-function.



Definition 26

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

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

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



Definition 27

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 :

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

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



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 .

Define

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

Lemma 28

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.

 $ax + b \not\equiv ay + b \pmod{p}$

If
$$x \neq y$$
 then $(x - y) \not\equiv 0 \pmod{p}$.

Multiplying with $a \not\equiv 0 \pmod{p}$ gives

$$a(x - y) \not\equiv 0 \pmod{p}$$

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 $t_y:=ay+b$.

This holds because we can compute a and b when given t_{χ} and t_{γ} :

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

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

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

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

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

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

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

Therefore, we can view the first step (before the $\operatorname{mod} n$ operation) as choosing a pair (t_x, t_y) , $t_x \neq t_y$ uniformly at
random.

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

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

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.

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

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 $t_x \mod n = h_1$ $(t_y \mod n = h_2)$ lies between $\lfloor \frac{p}{n} \rfloor$ and $\lceil \frac{p}{n} \rceil$.



Definition 29

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

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

The polynomial is defined by d+1 distinct points.

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

Let
$$A^{\ell} = \{h_{\bar{a}} \in \mathcal{H} \mid h_{\bar{a}}(x_i) = t_i \text{ for all } i \in \{1, \dots, \ell\}\}$$

Then

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

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

In order to obtain the cardinality of A^{ℓ} we choose our polynomial by fixing d+1 points.

We first fix the values for inputs x_1, \ldots, x_ℓ .

We have $\begin{vmatrix} \bullet & A^\ell & \text{denotes the set of hash-functions such that every } x_i \\ \text{hits its pre-defined position} \\ t_i. \end{aligned}$

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

 $(x_i) = t_i$). $\begin{cases} \bullet & B_i ext{ is the set of positions that} \\ & f_{ar{a}} ext{ can hit so that } h_{ar{a}} ext{ still hits} \\ & t_i. \end{cases}$

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 $h_{\bar{a}} \in A_{\ell}$.

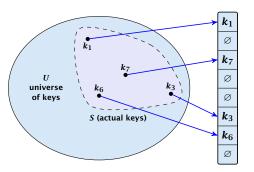
Therefore the probability of choosing $h_{ ilde{a}}$ from A_{ℓ} is only

$$\begin{split} & \frac{\lceil \frac{q}{n} \rceil^{\ell} \cdot q^{d-\ell+1}}{q^{d+1}} \leq \frac{(\frac{q+n}{n})^{\ell}}{q^{\ell}} \leq \left(\frac{q+n}{q}\right)^{\ell} \cdot \frac{1}{n^{\ell}} \\ & \leq \left(1 + \frac{1}{\ell}\right)^{\ell} \cdot \frac{1}{n^{\ell}} \leq \frac{e}{n^{\ell}} \ . \end{split}$$

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.



Let m = |S|. We could simply choose the hash-table size very large so that we don't get any collisions.

Using a universal hash-function the expected number of collisions is

$$E[\#Collisions] = \binom{m}{2} \cdot \frac{1}{n} .$$

If we choose $n=m^2$ the expected number of collisions is strictly less than $\frac{1}{2}$.

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 $\frac{1}{2}$.



297/530

We can find such a hash-function by a few trials.

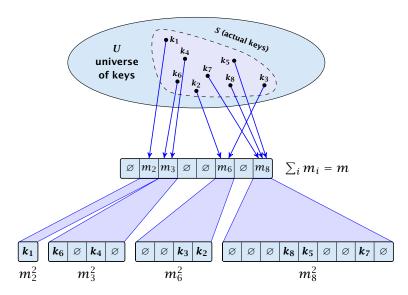
However, a hash-table size of $n = m^2$ is very very high.

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.



12. Jan. 2025 298/530





The total memory that is required by all hash-tables is $\mathcal{O}(\sum_j m_j^2)$. Note that m_j is a random variable.

$$E\left[\sum_{j} m_{j}^{2}\right] = E\left[2\sum_{j} {m_{j} \choose 2} + \sum_{j} m_{j}\right]$$
$$= 2E\left[\sum_{j} {m_{j} \choose 2}\right] + E\left[\sum_{j} m_{j}\right]$$

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!!!

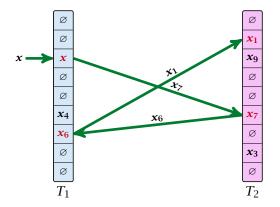


Goal:

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 $T_2[h_2(x)]$.
- A search clearly takes constant time if the above constraint is met.

Insert:



Algorithm 38 Cuckoo-Insert(x)

```
1: if T_1[h_1(x)] = x \vee T_2[h_2(x)] = x then return
2: steps ← 1
3: while steps ≤ 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}
     steps \leftarrow steps + 1
9: rehash() // change hash-functions; rehash everything
10: Cuckoo-Insert(x)
```

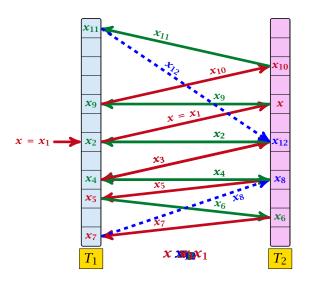
- We call one iteration through the while-loop a step of the algorithm.
- We call a sequence of iterations through the while-loop without the termination condition becoming true a phase of the algorithm.
- We say a phase is successful if it is not terminated by the maxstep-condition, but the while loop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

What is the expected time for an insert-operation?

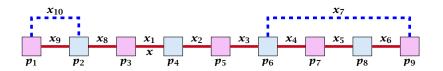
We first analyze the probability that we end-up in an infinite loop (that is then terminated after maxsteps steps).

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

Cuckoo Hashing: Insert



7.7 Hashing



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.
- 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.



12. Jan. 2025

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

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

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?

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

What is the probability that all keys in the cycle-structure of size s correctly map into their T_2 -cell?

This probability is at most $\frac{\mu}{n^s}$ since h_2 is a (μ, s) -independent hash-function.

These events are independent.



12. Jan. 2025

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

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

$$s^3 \cdot n^{s-1} \cdot m^{s-1}$$
.

- ► There are at most s^2 possibilities where to attach the forward and backward links.
- There are at most s possibilities to choose where to place key x.
- ▶ There are m^{s-1} possibilities to choose the keys apart from x.
- ▶ There are n^{s-1} possibilities to choose the cells.

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

$$\begin{split} \sum_{s=3}^{\infty} s^3 \cdot n^{s-1} \cdot m^{s-1} \cdot \frac{\mu^2}{n^{2s}} &= \frac{\mu^2}{nm} \sum_{s=3}^{\infty} s^3 \left(\frac{m}{n}\right)^s \\ &\leq \frac{\mu^2}{m^2} \sum_{s=3}^{\infty} s^3 \left(\frac{1}{1+\epsilon}\right)^s \leq \mathcal{O}\left(\frac{1}{m^2}\right) \ . \end{split}$$

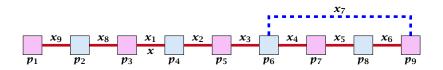
Here we used the fact that $(1 + \epsilon)m \le n$.

Hence,

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

313/530

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



Sequence of visited keys:

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

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

Lemma 30

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.

12. Jan. 2025 316/530

Taking $x_1 \rightarrow \cdots \rightarrow x_i$ twice, and $x_1 \rightarrow x_{i+1} \rightarrow \dots x_j$ once gives $2i + (j - i + 1) = i + j + 1 \ge p + 2$ keys. Hence, one of the sequences contains at least (p + 2)/3 keys.

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.

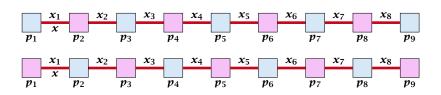
The sequence is of the form:

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

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

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

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

- 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 .



12. Jan. 2025

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.

Note that we count complete steps. A search that touches 2t or 2t + 1 keys takes t steps.

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

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

$$2 \cdot n^{s+1} \cdot m^{s-1} \cdot \frac{\mu^2}{n^{2s}}$$

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

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

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

12. Jan. 2025

We choose maxsteps $\geq 3\ell/2 + 1/2$. Then the probability that a phase terminates unsuccessfully without running into a cycle is at most

Pr[unsuccessful | no cycle]

$$\leq Pr[\,\exists \,\, \text{active path-structure of size at least} \,\, \frac{2\text{maxsteps}+2}{3}\,]$$

$$\leq \Pr[\exists \text{ active path-structure of size at least } \ell+1]$$

$$\leq \Pr[\exists \text{ active path-structure of size exactly } \ell+1]$$

$$\leq 2\mu^2 \Big(\frac{1}{1+\epsilon}\Big)^\ell \leq \frac{1}{m^2}$$

by choosing
$$\ell \ge \log\left(\frac{1}{2\mu^2m^2}\right)/\log\left(\frac{1}{1+\epsilon}\right) = \log\left(2\mu^2m^2\right)/\log\left(1+\epsilon\right)$$

This gives $\max steps = \Theta(\log m)$. Note that the existence of a path structure of ! size larger than s implies the existence of a path structure of size exactly s.

So far we estimated

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

and

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

Observe that

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

 $\geq c \cdot Pr[no cycle]$

for a suitable constant c > 0.

This is a very weak (and trivial) statement but still sufficient for our asymptotic analysis.

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

```
E[number of steps | phase successful]
= \sum_{t>1} \Pr[\text{search takes at least } t \text{ steps } | \text{ phase successful}]
```

We have

```
\begin{aligned} &\Pr[\mathsf{search} \; \mathsf{at} \; \mathsf{least} \; t \; \mathsf{steps} \; | \; \mathsf{successful}] \\ &= \Pr[\mathsf{search} \; \mathsf{at} \; \mathsf{least} \; t \; \mathsf{steps} \; \wedge \; \mathsf{successful}] / \Pr[\mathsf{successful}] \\ &\leq \frac{1}{c} \Pr[\mathsf{search} \; \mathsf{at} \; \mathsf{least} \; t \; \mathsf{steps} \; \wedge \; \mathsf{successful}] / \Pr[\mathsf{no} \; \mathsf{cycle}] \\ &\leq \frac{1}{c} \Pr[\mathsf{search} \; \mathsf{at} \; \mathsf{least} \; t \; \mathsf{steps} \; \wedge \; \mathsf{no} \; \mathsf{cycle}] / \Pr[\mathsf{no} \; \mathsf{cycle}] \\ &= \frac{1}{c} \Pr[\mathsf{search} \; \mathsf{at} \; \mathsf{least} \; t \; \mathsf{steps} \; | \; \mathsf{no} \; \mathsf{cycle}] \; . \end{aligned}
```

Hence,

E[number of steps | phase successful]

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

$$\leq \frac{1}{c} \sum_{t \geq 1} 2\mu^2 \left(\frac{1}{1+\epsilon}\right)^{(2t-1)/3} = \frac{1}{c} \sum_{t \geq 0} 2\mu^2 \left(\frac{1}{1+\epsilon}\right)^{(2(t+1)-1)/3}$$

$$= \frac{2\mu^2}{c(1+\epsilon)^{1/3}} \sum_{t \geq 0} \left(\frac{1}{(1+\epsilon)^{2/3}}\right)^t = \mathcal{O}(1) .$$

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).

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).

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

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



12. Jan. 2025

Formal Proof

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):

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

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

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

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):

$$\begin{split} \mathbf{E}[X_i^s] &= \mathbf{E}[\mathsf{steps} \mid \mathsf{phase} \; \mathsf{successful}] \cdot \Pr[\mathsf{phase} \; \mathsf{sucessful}] \\ &+ \mathsf{maxsteps} \cdot \Pr[\mathsf{not} \; \mathsf{sucessful}] = \mathcal{O}(1) \;\;. \end{split}$$

The expected cost for all rehashes is

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

Note that Z_i is independent of X_j^s , $j \ge i$ (however, it is not independent of X_j^s , j < i). Hence,

$$\begin{split} \mathbf{E}\left[\sum_{i}\sum_{s}Z_{i}X_{s}^{i}\right] &= \sum_{i}\sum_{s}\mathbf{E}[Z_{i}]\cdot\mathbf{E}[X_{s}^{i}] \\ &\leq \mathcal{O}(m)\cdot\sum_{i}p^{i} \\ &\leq \mathcal{O}(m)\cdot\frac{p}{1-p} \\ &= \mathcal{O}(1) \ . \end{split}$$

What kind of hash-functions do we need?

Since maxsteps is $\Theta(\log m)$ the largest size of a path-structure or cycle-structure contains just $\Theta(\log m)$ different keys.

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

12. Jan. 2025

How do we make sure that $n \ge (1 + \epsilon)m$?

- ▶ Let $\alpha := 1/(1 + \epsilon)$.
- Keep track of the number of elements in the table. When $m \ge \alpha n$ we double n and do a complete re-hash (table-expand).
- Whenever m drops below $\alpha n/4$ we divide n by 2 and do a rehash (table-shrink).
- Note that right after a change in table-size we have $m = \alpha n/2$. In order for a table-expand to occur at least $\alpha n/2$ insertions are required. Similar, for a table-shrink at least $\alpha n/4$ deletions must occur.
- Therefore we can amortize the rehash cost after a change in table-size against the cost for insertions and deletions.

12. Jan. 2025 329/530

Lemma 31

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

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 $m \le (1+\epsilon)n$.



Hashing

Bibliography

[MS08] Kurt Mehlhorn, Peter Sanders: Algorithms and Data Structures — The Basic Toolbox, Springer, 2008

[CLRS90] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein: Introduction to algorithms (3rd ed.), MIT Press and McGraw-Hill, 2009

Chapter 4 of [MS08] contains a detailed description about Hashing with Linear Probing and Hashing with Chaining. Also the Perfect Hashing scheme can be found there.

The analysis of Hashing with Chaining under the assumption of uniform hashing can be found in Chapter 11.2 of [CLRS90]. Chapter 11.3.3 describes Universal Hashing. Collision resolution with Open Addressing is described in Chapter 11.4. Chapter 11.5 describes the Perfect Hashing scheme.

Reference for Cuckoo Hashing???

12. Jan. 2025

8 Priority Queues

A Priority Queue S is a dynamic set data structure that supports the following operations:

- ▶ *S.* build($x_1, ..., x_n$): Creates a data-structure that contains just the elements $x_1, ..., x_n$.
- \triangleright S. insert(x): Adds element x to the data-structure.
- ▶ element *S*. minimum(): Returns an element $x \in S$ with minimum key-value key[x].
- element S. delete-min(): Deletes the element with minimum key-value from S and returns it.
- **boolean** *S.* **is-empty**(): Returns true if the data-structure is empty and false otherwise.

Sometimes we also have

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



331/530

8 Priority Queues

An addressable Priority Queue also supports:

- ► handle S. insert(x): Adds element x to the data-structure, and returns a handle to the object for future reference.
- **S.** delete(h): Deletes element specified through handle h.
- 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 39 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. delete-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 40 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
 4: for all v \in V \setminus \{s\} do
 5: v \cdot \ker \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. delete-min();
10: for all x \in V s.t. \{v, x\} \in E do
11:
                if x. key > d(v, x) then
12:
                      S.decrease-key(h_x,d(v,x));
                      x. key \leftarrow d(v, x);
13:
14:
                      x. pred \leftarrow v:
```



Analysis of Dijkstra and Prim

Both algorithms require:

- 1 build() operation
- ightharpoonup |V| insert() operations
- ▶ |V| delete-min() operations
- ▶ |V| is-empty() operations
- \triangleright |E| decrease-key() operations

How good a running time can we obtain?

8 Priority Queues

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

Note that most applications use build() only to create an empty heap which then costs time 1.

** The standard version of binary heaps is not addressable. Hence, it does not support a delete.

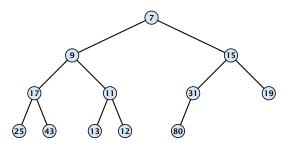
8 Priority Queues

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

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

8.1 Binary Heaps

- Nearly complete binary tree; only the last level is not full, and this one is filled from left to right.
- Heap property: A node's key is not larger than the key of one of its children.



Binary Heaps

Operations:

- **minimum():** return the root-element. Time O(1).
- **is-empty**(): check whether root-pointer is null. Time $\mathcal{O}(1)$.

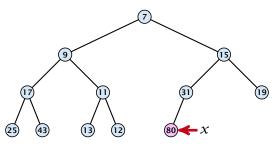
8.1 Binary Heaps

Maintain a pointer to the last element x.

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

go up until the last edge used was a right edge. go left; go right until you reach a leaf

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



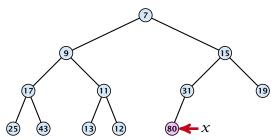
8.1 Binary Heaps

Maintain a pointer to the last element x.

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

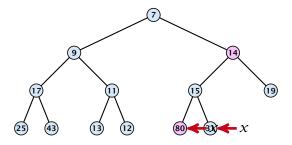
go up until the last edge used was a left edge. go right; go left until you reach a null-pointer.

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



Insert

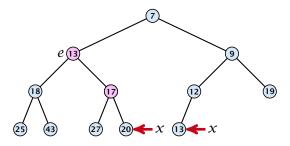
- 1. Insert element at successor of x.
- 2. Exchange with parent until heap property is fulfilled.



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.

Delete

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



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 $O(\log n)$.

Binary Heaps

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 $\mathcal{O}(n)$.

Binary Heaps

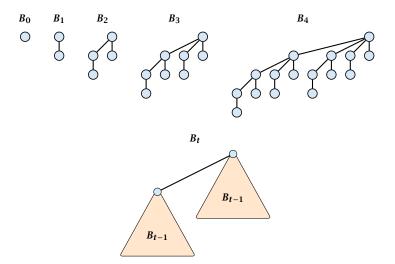
The standard implementation of binary heaps is via arrays. Let A[0,...,n-1] be an array

- ▶ 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.

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

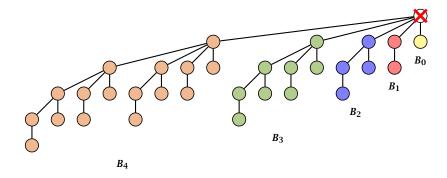
The resulting binary heap is not addressable. The elements don't maintain their positions and therefore there are no stable handles.

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

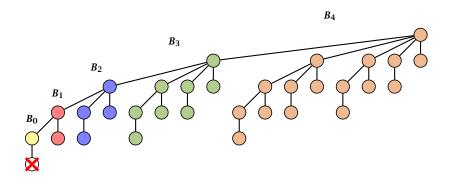


Properties of Binomial Trees

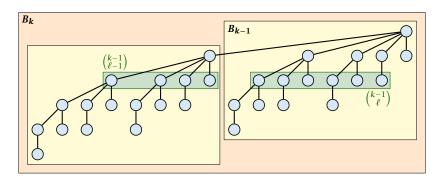
- \triangleright B_k has 2^k nodes.
- $ightharpoonup B_k$ has height k.
- ▶ The root of B_k has degree k.
- ▶ B_k has $\binom{k}{\ell}$ nodes on level ℓ .
- ▶ Deleting the root of B_k gives trees $B_0, B_1, ..., B_{k-1}$.



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



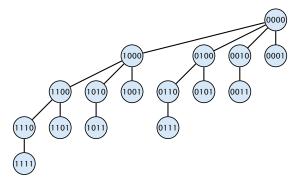
Deleting the leaf furthest from the root (in B_5) leaves a path that connects the roots of sub-trees B_4 , B_3 , B_2 , B_1 , and B_0 .



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

$$\binom{k-1}{\ell-1}+\binom{k-1}{\ell}=\binom{k}{\ell}$$





The binomial tree B_k is a sub-graph of the hypercube H_k .

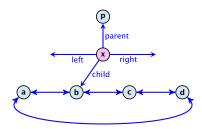
The parent of a node with label $b_k, ..., b_1$ is obtained by setting the least significant 1-bit to 0.

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



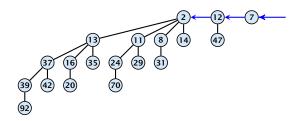
How do we implement trees with non-constant degree?

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



- 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*.

Binomial Heap



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

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

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.

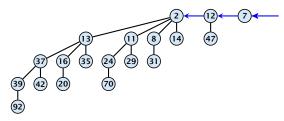
Let B_{k_1} , B_{k_2} , B_{k_3} , $k_i < k_{i+1}$ denote the binomial trees in the collection and recall that every tree may be contained at most once.

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.

Binomial Heap

Properties of a heap with n keys:

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



Binomial Heap: Merge

The merge-operation is instrumental for binomial heaps.

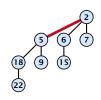
A merge is easy if we have two heaps with different binomial trees. We can simply merge the tree-lists.

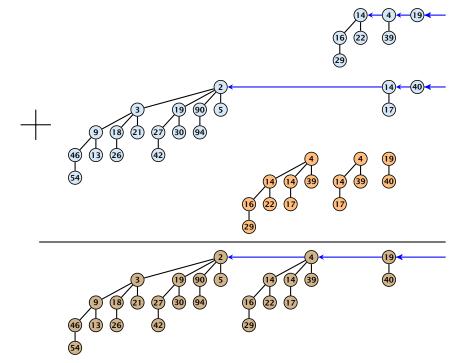
Note that we do not just do a concatenation as we want to keep the trees in the list sorted according to size.

Otherwise, we cannot do this because the merged heap is not allowed to contain two trees of the same order.

Merging two trees of the same size: Add the tree with larger root-value as a child to the other tree.

For more trees the technique is analogous to binary addition.





S_1 . merge(S_2):

- Analogous to binary addition.
- Time is proportional to the number of trees in both heaps.
- ▶ Time: $O(\log n)$.

All other operations can be reduced to merge().

S. insert(x):

- Create a new heap S' that contains just the element x.
- ightharpoonup Execute S. merge(S').
- ▶ Time: $O(\log n)$.

S. minimum():

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

S. delete-min():

- Find the minimum key-value among all roots.
- Remove the corresponding tree T_{\min} from the heap.
- Create a new heap S' that contains the trees obtained from T_{\min} after deleting the root (note that these are just $\mathcal{O}(\log n)$ trees).
- \triangleright Compute S. merge(S').
- ightharpoonup Time: $\mathcal{O}(\log n)$.

S. decrease-key(handle h):

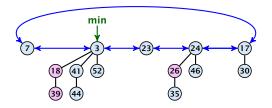
- lacktriangle Decrease the key of the element pointed to by h.
- Bubble the element up in the tree until the heap property is fulfilled.
- ▶ Time: $O(\log n)$ since the trees have height $O(\log n)$.

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.

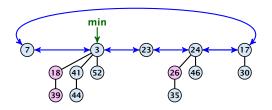


Additional implementation details:

- 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 \boldsymbol{c} to denote the amount of work that a unit of potential can pay for.

S. minimum()

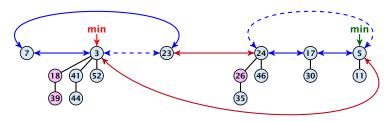
- Access through the min-pointer.
- ightharpoonup Actual cost $\mathcal{O}(1)$.
- No change in potential.
- ▶ Amortized cost $\mathcal{O}(1)$.

S. merge(S')

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

In the figure below the dashed edges are replaced by red edges.

 The minimum of the left heap becomes the new minimum of the merged heap.



Running time:

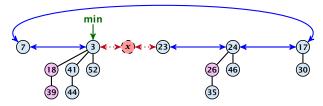
- ightharpoonup Actual cost $\mathcal{O}(1)$.
- No change in potential.
- ▶ Hence, amortized cost is $\mathcal{O}(1)$.



x is inserted next to the min-pointer as this is our entry point into the root-list.

S. insert(x)

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



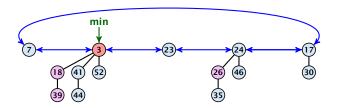
Running time:

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

 $D(\min)$ is the number of children of the node that stores the minimum.

S. delete-min(x)

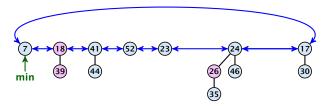
- Delete minimum; add child-trees to heap; time: $D(\min) \cdot \mathcal{O}(1)$.
- Update min-pointer; time: $(t + D(\min)) \cdot \mathcal{O}(1)$.



 $D(\min)$ is the number of children of the node that stores the minimum.

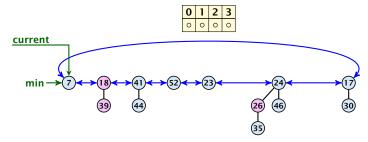
S. delete-min(x)

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



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

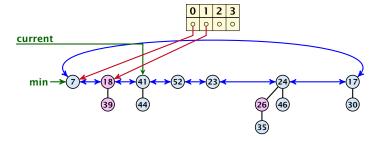
Consolidate:

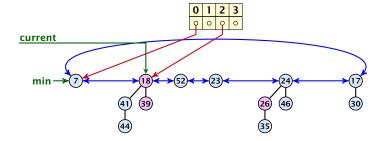


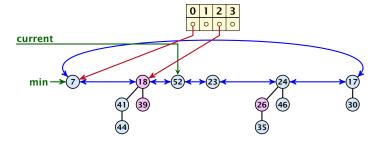
During the consolidation we traverse the root list. Whenever we discover two trees that have the same degree we merge these trees. In order to efficiently check whether two trees have the same degree, we use an array that contains for every degree value d a pointer to a tree left of the current pointer whose root has degree d (if such a tree exist).

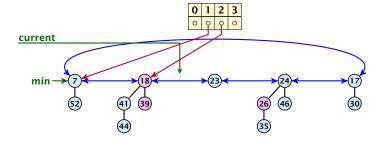
12. Jan. 2025

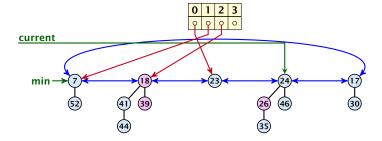
Consolidate:

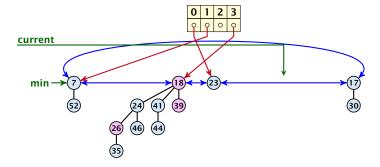


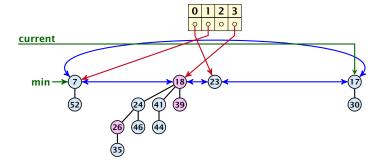


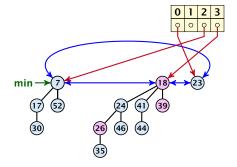












t and t' denote the number of trees before and after the delete-min() operation, respectively. D_n is an upper bound on the degree (i.e., number of children) of a tree node.

Actual cost for delete-min()

- At most $D_n + t$ elements in root-list before consolidate.
- Actual cost for a delete-min is at most $\mathcal{O}(1) \cdot (D_n + t)$. Hence, there exists c_1 s.t. actual cost is at most $c_1 \cdot (D_n + t)$.

Amortized cost for delete-min()

- $ightharpoonup t' \leq D_n + 1$ as degrees are different after consolidating.
- ► Therefore $\Delta \Phi \leq D_n + 1 t$;
- We can pay $c \cdot (t D_n 1)$ from the potential decrease.
- The amortized cost is

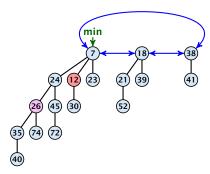
$$c_1 \cdot (D_n + t) - c \cdot (t - D_n - 1)$$

 $\leq (c_1 + c)D_n + (c_1 - c)t + c \leq 2c(D_n + 1) \leq \mathcal{O}(D_n)$

for $c \ge c_1$.

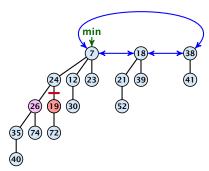
If the input trees of the consolidation procedure are binomial trees (for example only singleton vertices) then the output will be a set of distinct binomial trees, and, hence, the Fibonacci heap will be (more or less) a Binomial heap right after the consolidation.

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

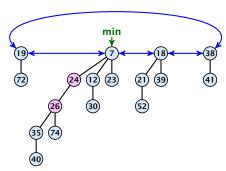
▶ Just decrease the key-value of element referenced by h. Nothing else to do.



Case 2: heap-property is violated, but parent is not marked

- Decrease key-value of element x reference by h.
- If the heap-property is violated, cut the parent edge of x, and make x into a root.
- Adjust min-pointers, if necessary.
- \blacktriangleright Mark the (previous) parent of x (unless it's a root).



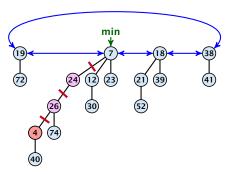


Case 2: heap-property is violated, but parent is not marked

- Decrease key-value of element x reference by h.
- If the heap-property is violated, cut the parent edge of x, and make x into a root.
- Adjust min-pointers, if necessary.
- \blacktriangleright Mark the (previous) parent of x (unless it's a root).



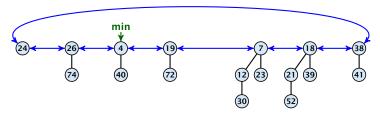
378/530



Case 3: heap-property is violated, and parent is marked

- ▶ Decrease key-value of element x reference by h.
- Let the parent edge of x, and make x into a root.
- Adjust min-pointers, if necessary.
- Continue cutting the parent until you arrive at an unmarked node.





Case 3: heap-property is violated, and parent is marked

- ▶ Decrease key-value of element x reference by h.
- Let the parent edge of x, and make x into a root.
- Adjust min-pointers, if necessary.
- Continue cutting the parent until you arrive at an unmarked node.



Case 3: heap-property is violated, and parent is marked

- ightharpoonup Decrease key-value of element x reference by h.
- Cut the parent edge of x, and make x into a root.
- Adjust min-pointers, if necessary.
- Execute the following:

```
p \leftarrow \operatorname{parent}[x]; marked; the some shift is made while (p is marked) pp \leftarrow \operatorname{parent}[p]; cut of p; make it into a root; unmark it; p \leftarrow pp;
```

if p is unmarked and not a root mark it;

Marking a node can be viewed as a first step towards becoming a root. The first time x loses a child it is marked; the second time it loses a child it is made into a root.

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 .

Amortized cost:

- $t' = t + \ell$, 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

$$c_2(\ell+1)+c(4-\ell) \leq (c_2-c)\ell+4c+c_2=\mathcal{O}(1)$$
, m and m' : number of marked nodes before if $c \geq c_2$.

t and t': number of trees before and after operation.
m and m': number of marked nodes before and after operation.



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 32

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

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

The marking process is very important for the proof of this lemma. It ensures that a node can have lost at most one child since the last time it became a non-root node. When losing a first child the node gets marked; when losing the second child it is cut from the parent and made into a root.

Proof

- When y_i was linked to x, at least y_1, \ldots, y_{i-1} were already linked to x.
- ▶ Hence, at this time $degree(x) \ge i 1$, and therefore also $degree(y_i) \ge i 1$ as the algorithm links nodes of equal degree only.
- 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.
- $ightharpoonup s_k$ monotonically increases with k
- $ightharpoonup s_0 = 1 \text{ and } s_1 = 2.$

Let x be a degree k node of size s_k and let y_1, \ldots, y_k be its children.

$$s_k = 2 + \sum_{i=2}^k \operatorname{size}(y_i)$$

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

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

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

Definition 33

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



Priority Queues

Bibliography

[CLRS90] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein: Introduction to algorithms (3rd ed.), MIT Press and McGraw-Hill, 2009

[MS08] Kurt Mehlhorn, Peter Sanders: Algorithms and Data Structures — The Basic Toolbox, Springer, 2008

Binary heaps are covered in [CLRS90] in combination with the heapsort algorithm in Chapter 6. Fibonacci heaps are covered in detail in Chapter 19. Problem 19-2 in this chapter introduces Binomial heaps.

Chapter 6 in [MS08] covers Priority Queues. Chapter 6.2.2 discusses Fibonacci heaps. Binomial heaps are dealt with in Exercise 6.11.



9 Union Find

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

- ▶ **P.** makeset(x): Given an element x, adds x to the data-structure and creates a singleton set that contains only this element. Returns a locator/handle for x in the data-structure.
- ▶ \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.

12. Jan. 2025

9 Union Find

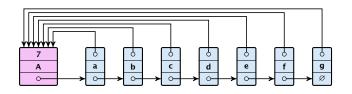
Applications:

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

9 Union Find

Algorithm 41 Kruskal-MST(G = (V, E), w)1: $A \leftarrow \emptyset$; 2: **for all** $v \in V$ **do**3: $v. \sec \leftarrow \mathcal{P}. \text{makeset}(v. \text{label})$ 4: sort edges in non-decreasing order of weight w5: **for all** $(u, v) \in E$ in non-decreasing order **do**6: **if** $\mathcal{P}. \text{find}(u. \sec) \neq \mathcal{P}. \text{find}(v. \sec)$ **then**7: $A \leftarrow A \cup \{(u, v)\}$ 8: $\mathcal{P}. \text{union}(u. \sec, v. \sec)$

- ► The elements of a set are stored in a list; each node has a backward pointer to the head.
- ► The head of the list contains the identifier for the set and a field that stores the size of the set.



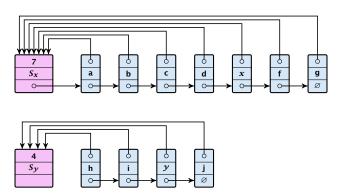
- ightharpoonup makeset(x) can be performed in constant time.
- find(x) can be performed in constant time.



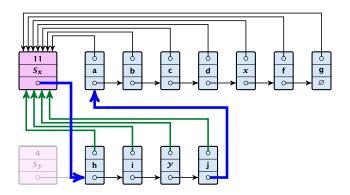
union(x, y)

- ▶ Determine sets S_X and S_Y .
- ► Traverse the smaller list (say S_y), and change all backward pointers to the head of list S_x .
- ▶ Insert list S_{γ} at the head of S_{χ} .
- ▶ Adjust the size-field of list S_x .
- ► Time: $\min\{|S_x|, |S_y|\}$.

391/530









12. Jan. 2025 **392/530**

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 34

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

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

The Accounting Method for Amortized Time Bounds

- There is a bank account for every element in the data structure.
- Initially the balance on all accounts is zero.
- Whenever for an operation the amortized time bound exceeds the actual cost, the difference is credited to some bank accounts of elements involved.
- Whenever for an operation the actual cost exceeds the amortized time bound, the difference is charged to bank accounts of some of the elements involved.
- If we can find a charging scheme that guarantees that balances always stay positive the amortized time bounds are proven.



- For an operation whose actual cost exceeds the amortized cost we charge the excess to the elements involved.
- In total we will charge at most $O(\log n)$ to an element (regardless of the request sequence).
- For each element a makeset operation occurs as the first operation involving this element.
- ▶ We inflate the amortized cost of the makeset-operation to $\Theta(\log n)$, i.e., at this point we fill the bank account of the element to $\Theta(\log n)$.
- Later operations charge the account but the balance never drops below zero.



12. Jan. 2025 396/530

makeset(x): The actual cost is O(1). Due to the cost inflation the amortized cost is $O(\log n)$.

find(x): For this operation we define the amortized cost and the actual cost to be the same. Hence, this operation does not change any accounts. Cost: $\mathcal{O}(1)$.

union(x, y):

- ▶ If $S_x = S_y$ the cost is constant; no bank accounts change.
- ▶ Otw. the actual cost is $\mathcal{O}(\min\{|S_x|, |S_y|\})$.
- Assume wlog. that S_x is the smaller set; let c denote the hidden constant, i.e., the actual cost is at most $c \cdot |S_x|$.
- ▶ Charge c to every element in set S_x .



Lemma 35

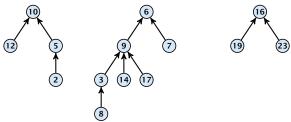
An element is charged at most $\lfloor \log_2 n \rfloor$ times, where n is the total number of elements in the set system.

Proof.

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



- Maintain nodes of a set in a tree.
- 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.
- Example:



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

399/530

makeset(x)

- Create a singleton tree. Return pointer to the root.
- ightharpoonup Time: $\mathcal{O}(1)$.

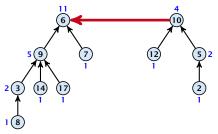
find(x)

- 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.

To support union we store the size of a tree in its root.

union(x, y)

- ▶ Perform $a \leftarrow \text{find}(x)$; $b \leftarrow \text{find}(y)$. Then: link(a, b).
- \blacktriangleright link(a, b) attaches the smaller tree as the child of the larger.
- In addition it updates the size-field of the new root.



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

Lemma 36

The running time (non-amortized!!!) for find(x) is $O(\log n)$.

Proof.

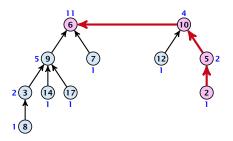
- ▶ When we attach a tree with root c to become a child of a tree with root p, then $\operatorname{size}(p) \ge 2\operatorname{size}(c)$, where size denotes the value of the size-field right after the operation.
- After that the value of size(c) stays fixed, while the value of size(p) may still increase.
- ► Hence, at any point in time a tree fulfills $size(p) \ge 2 \, size(c)$, for any pair of nodes (p,c), where p is a parent of c.



Path Compression

find(x):

- Go upward until you find the root.
- Re-attach all visited nodes as children of the root.
- Speeds up successive find-operations.

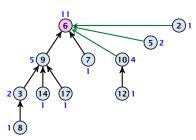


Note that the size-fields now only give an upper bound on the size of a sub-tree.

Path Compression

find(x):

- Go upward until you find the root.
- Re-attach all visited nodes as children of the root.
- Speeds up successive find-operations.



One could change the algorithm to update the size-fields. This could be done without asymptotically affecting the running time.

However, the only size-field that is actually required is the field at the root, which is always correct.

We will only use the other size-fields for the proof of Theorem 39.

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

Path Compression

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

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

Definitions:

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).

Note that this is the same as the size of ν 's subtree in the case that there are no find-operations.

- $ightharpoonup rank(v) = \lfloor \log(\text{size}(v)) \rfloor.$
- \Rightarrow size $(v) \ge 2^{\operatorname{rank}(v)}$.

Lemma 37

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



Lemma 38

There are at most $n/2^s$ nodes of rank s.

Proof.

- Let's say a node v sees node x if v is in x's sub-tree at the time that x becomes a child.
- 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.

We define

and

$$\log^*(n) := \min\{i \mid \text{tow}(i) \ge n\} .$$

Theorem 39

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) : $\mathcal{O}(\log^*(n))$

In the following we assume $n \ge 2$.

rank-group:

- ▶ A node with rank rank(v) is in rank group $log^*(rank(v))$.
- ▶ The rank-group g = 0 contains only nodes with rank 0 or rank 1.
- ▶ A rank group $g \ge 1$ contains ranks tow(g-1) + 1, ..., tow(g).
- ► 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 $\log^* n$.



Accounting Scheme:

- create an account for every find-operation
- lacktriangle create an account for every node v

The cost for a find-operation is equal to the length of the path traversed. We charge the cost for going from v to parent[v] as follows:

- If parent[v] is the root we charge the cost to the find-account.
- If the group-number of rank(v) is the same as that of rank(parent[v]) (before starting path compression) we charge the cost to the node-account of v.
- Otherwise we charge the cost to the find-account.

12. Jan. 2025

409/530

Observations:

- A find-account is charged at most $\log^*(n)$ times (once for the root and at most $\log^*(n) 1$ times when increasing the rank-group).
- ightharpoonup 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 $tow(g) tow(g-1) 1 \le tow(g)$.

What is the total charge made to nodes?

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

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

$$= \frac{n}{2^{\text{tow}(g-1)+1}} \sum_{s=0}^{\infty} \frac{1}{2^s} = \frac{n}{2^{\text{tow}(g-1)+1}} \cdot 2$$

$$= \frac{n}{2^{\text{tow}(g-1)}} = \frac{n}{\text{tow}(g)}.$$

Hence,

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

Without loss of generality we can assume that all makeset-operations occur at the start.

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).

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

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

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

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

Union Find

Bibliography

- [CLRS90a] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest: Introduction to Algorithms (1st ed.), MIT Press and McGraw-Hill. 1990
- [CLRS90b] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein: Introduction to Algorithms (2nd ed.), MIT Press and McGraw-Hill, 2001
- [CLRS90c] Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein: Introduction to Algorithms (3rd ed.), MIT Press and McGraw-Hill. 2009
- [AHU74] Alfred V. Aho, John E. Hopcroft, Jeffrey D. Ullman: The Design and Analysis of Computer Algorithms, Addison-Wesley, 1974

Union find data structures are discussed in Chapter 21 of [CLRS90b] and [CLRS90c] and in Chapter 22 of [CLRS90a]. The analysis of union by rank with path compression can be found in [CLRS90a] but neither in [CLRS90b] in nor in [CLRS90c]. The latter books contains a more involved analysis that gives a better bound than $\mathcal{O}(\log^* n)$.

A description of the $\mathcal{O}(\log^*)$ -bound can also be found in Chapter 4.8 of [AHU74].



Part IV

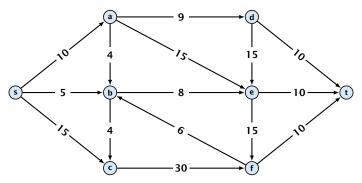
Flows and Cuts

The following slides are partially based on slides by Kevin Wayne.

10 Introduction

Flow Network

- directed graph G = (V, E); edge capacities c(e)
- two special nodes: source s; target t;
- no edges entering s or leaving t;
- at least for now: no parallel edges;



Cuts

Definition 40

An (s, t)-cut in the graph G is given by a set $A \subset V$ with $s \in A$ and $t \in V \setminus A$.

Definition 41

The capacity of a cut A is defined as

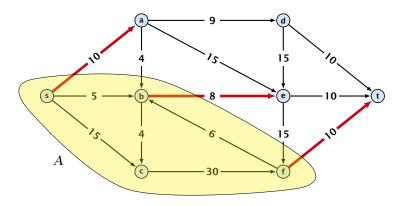
$$cap(A, V \setminus A) := \sum_{e \in out(A)} c(e) ,$$

where $\operatorname{out}(A)$ denotes the set of edges of the form $A \times V \setminus A$ (i.e. edges leaving A).

Minimum Cut Problem: Find an (s, t)-cut with minimum capacity.

Cuts

Example 42



The capacity of the cut is $cap(A, V \setminus A) = 28$.



Definition 43

An (s,t)-flow is a function $f: E \mapsto \mathbb{R}^+$ that satisfies

1. For each edge e

$$0 \le f(e) \le c(e)$$
.

(capacity constraints)

2. For each $v \in V \setminus \{s, t\}$

$$\sum_{e \in \text{out}(v)} f(e) = \sum_{e \in \text{into}(v)} f(e) .$$

(flow conservation constraints)

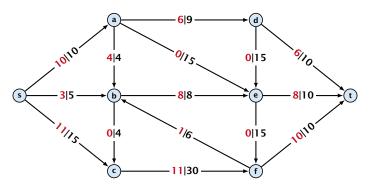
Definition 44

The value of an (s, t)-flow f is defined as

$$val(f) = \sum_{e \in out(s)} f(e)$$
.

Maximum Flow Problem: Find an (s, t)-flow with maximum value.

Example 45



The value of the flow is val(f) = 24.

Lemma 46 (Flow value lemma)

Let f be a flow, and let $A \subseteq V$ be an (s,t)-cut. Then the net-flow across the cut is equal to the amount of flow leaving s, i.e.,

$$\operatorname{val}(f) = \sum_{e \in \operatorname{out}(A)} f(e) - \sum_{e \in \operatorname{into}(A)} f(e)$$
.

Proof.

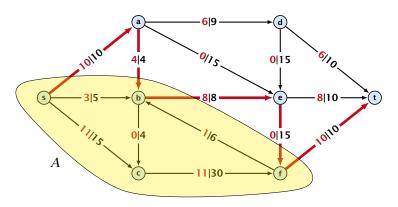
$$val(f) = \sum_{e \in out(s)} f(e)$$

$$= \sum_{e \in out(s)} f(e) + \sum_{v \in A \setminus \{s\}} \left(\sum_{e \in out(v)} f(e) - \sum_{e \in in(v)} f(e) \right)$$

$$= \sum_{e \in out(A)} f(e) - \sum_{e \in into(A)} f(e)$$

The last equality holds since every edge with both end-points in A contributes negatively as well as positively to the sum in Line 2. The only edges whose contribution doesn't cancel out are edges leaving or entering A.

Example 47



The net-flow across the cut is val(f) = 24.

10 Introduction

Corollary 48

Let f be an (s,t)-flow and let A be an (s,t)-cut, such that

$$\operatorname{val}(f) = \operatorname{cap}(A, V \setminus A).$$

Then f is a maximum flow.

Proof.

Suppose that there is a flow f' with larger value. Then

$$cap(A, V \setminus A) < val(f')$$

$$= \sum_{e \in out(A)} f'(e) - \sum_{e \in into(A)} f'(e)$$

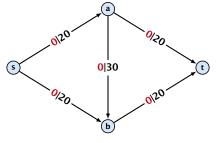
$$\leq \sum_{e \in out(A)} f'(e)$$

$$\leq cap(A, V \setminus A)$$

11 Augmenting Path Algorithms

Greedy-algorithm:

- start with f(e) = 0 everywhere
- find an s-t path with f(e) < c(e) on every edge
- augment flow along the path
- repeat as long as possible

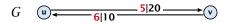


fflow weature: 200

The Residual Graph

From the graph G = (V, E, c) and the current flow f we construct an auxiliary graph $G_f = (V, E_f, c_f)$ (the residual graph):

- Suppose the original graph has edges $e_1 = (u, v)$, and $e_2 = (v, u)$ between u and v.
- ▶ G_f has edge e_1' with capacity $\max\{0, c(e_1) f(e_1) + f(e_2)\}$ and e_2' with with capacity $\max\{0, c(e_2) f(e_2) + f(e_1)\}$.



$$G_f = 0$$
 $\longrightarrow 0$ $\longrightarrow 0$

Augmenting Path Algorithm

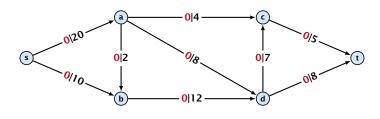
Definition 49

An augmenting path with respect to flow f, is a path from s to t in the auxiliary graph G_f that contains only edges with non-zero capacity.

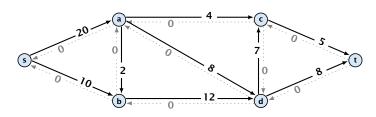
Algorithm 1 FordFulkerson(G = (V, E, c))

- 1: Initialize $f(e) \leftarrow 0$ for all edges.
- 2: **while** \exists augmenting path p in G_f **do**
- 3: augment as much flow along p as possible.

Augmenting Paths



flow value: 0



Augmenting Path Algorithm

Theorem 50

A flow f is a maximum flow **iff** there are no augmenting paths.

Theorem 51

The value of a maximum flow is equal to the value of a minimum cut.

Proof.

Let f be a flow. The following are equivalent:

- **1.** There exists a cut A such that $val(f) = cap(A, V \setminus A)$.
- 2. Flow f is a maximum flow.
- 3. There is no augmenting path w.r.t. f.



Augmenting Path Algorithm

$$1. \Rightarrow 2.$$

This we already showed.

$$2. \Rightarrow 3.$$

If there were an augmenting path, we could improve the flow. Contradiction.

- $3. \Rightarrow 1.$
 - Let f be a flow with no augmenting paths.
 - Let A be the set of vertices reachable from s in the residual graph along non-zero capacity edges.
 - ▶ Since there is no augmenting path we have $s \in A$ and $t \notin A$.

Augmenting Path Algorithm

$$val(f) = \sum_{e \in out(A)} f(e) - \sum_{e \in into(A)} f(e)$$
$$= \sum_{e \in out(A)} c(e)$$
$$= cap(A, V \setminus A)$$

This finishes the proof.

Here the first equality uses the flow value lemma, and the second exploits the fact that the flow along incoming edges must be 0 as the residual graph does not have edges leaving A.

Analysis

Assumption:

All capacities are integers between 1 and C.

Invariant:

Every flow value $f(\emph{e})$ and every residual capacity $\emph{c}_f(\emph{e})$ remains integral troughout the algorithm.

Lemma 52

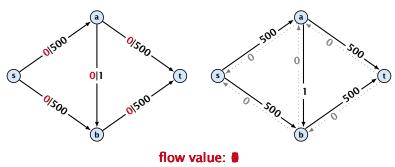
The algorithm terminates in at most $val(f^*) \le nC$ iterations, where f^* denotes the maximum flow. Each iteration can be implemented in time $\mathcal{O}(m)$. This gives a total running time of $\mathcal{O}(nmC)$.

Theorem 53

If all capacities are integers, then there exists a maximum flow for which every flow value f(e) is integral.

A Bad Input

Problem: The running time may not be polynomial

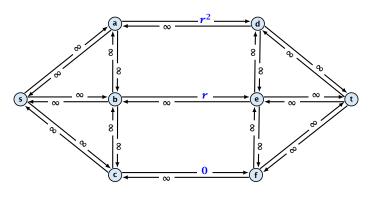


Question:

Can we tweak the algorithm so that the running time is polynomial in the input length?

A Pathological Input

Let
$$r = \frac{1}{2}(\sqrt{5} - 1)$$
. Then $r^{n+2} = r^n - r^{n+1}$.



flo**flottheetice:** 19^{2} 14^{3} r^{4}

Running time may be infinite!!!

How to choose augmenting paths?

- We need to find paths efficiently.
- We want to guarantee a small number of iterations.

Several possibilities:

- Choose path with maximum bottleneck capacity.
- Choose path with sufficiently large bottleneck capacity.
- Choose the shortest augmenting path.

Overview: Shortest Augmenting Paths

Lemma 54

The length of the shortest augmenting path never decreases.

Lemma 55

After at most $\mathcal{O}(m)$ augmentations, the length of the shortest augmenting path strictly increases.

Overview: Shortest Augmenting Paths

These two lemmas give the following theorem:

Theorem 56

The shortest augmenting path algorithm performs at most $\mathcal{O}(mn)$ augmentations. This gives a running time of $\mathcal{O}(m^2n)$.

Proof.

- We can find the shortest augmenting paths in time $\mathcal{O}(m)$ via BFS.
- O(m) augmentations for paths of exactly k < n edges.



Define the level $\ell(v)$ of a node as the length of the shortest s-v path in G_f (along non-zero edges).

Let L_G denote the subgraph of the residual graph G_f that contains only those edges (u, v) with $\ell(v) = \ell(u) + 1$.

A path P is a shortest s-u path in G_f iff it is an s-u path in L_G .





In the following we assume that the residual graph \mathcal{G}_f does not contain zero capacity edges.

This means, we construct it in the usual sense and then delete edges of zero capacity.

First Lemma:

The length of the shortest augmenting path never decreases.

After an augmentation G_f changes as follows:

- Bottleneck edges on the chosen path are deleted.
- Back edges are added to all edges that don't have back edges so far.

These changes cannot decrease the distance between s and t.

Second Lemma: After at most m augmentations the length of the shortest augmenting path strictly increases.

Let M denote the set of edges in graph L_G at the beginning of a round when the distance between s and t is k.

An s-t path in G_f that uses edges not in M has length larger than k, even when using edges added to G_f during the round.

In each augmentation an edge is deleted from M.

non-shortest path.

Theorem 57

The shortest augmenting path algorithm performs at most $\mathcal{O}(mn)$ augmentations. Each augmentation can be performed in time $\mathcal{O}(m)$.

Theorem 58 (without proof)

There exist networks with $m = \Theta(n^2)$ that require $\Omega(mn)$ augmentations, when we restrict ourselves to only augment along shortest augmenting paths.

Note:

There always exists a set of m augmentations that gives a maximum flow (why?).

When sticking to shortest augmenting paths we cannot improve (asymptotically) on the number of augmentations.

However, we can improve the running time to $\mathcal{O}(mn^2)$ by improving the running time for finding an augmenting path (currently we assume $\mathcal{O}(m)$ per augmentation for this).

We maintain a subset M of the edges of G_f with the guarantee that a shortest s-t path using only edges from M is a shortest augmenting path.

With each augmentation some edges are deleted from M.

When M does not contain an s-t path anymore the distance between s and t strictly increases.

Note that ${\cal M}$ is not the set of edges of the level graph but a subset of level-graph edges.

Suppose that the initial distance between s and t in G_f is k.

M is initialized as the level graph L_G .

Perform a DFS search to find a path from s to t using edges from M.

Either you find t after at most n steps, or you end at a node v that does not have any outgoing edges.

You can delete incoming edges of v from M.

Analysis

Let a phase of the algorithm be defined by the time between two augmentations during which the distance between s and t strictly increases.

Initializing M for the phase takes time $\mathcal{O}(m)$.

The total cost for searching for augmenting paths during a phase is at most $\mathcal{O}(mn)$, since every search (successful (i.e., reaching t) or unsuccessful) decreases the number of edges in M and takes time $\mathcal{O}(n)$.

The total cost for performing an augmentation during a phase is only $\mathcal{O}(n)$. For every edge in the augmenting path one has to update the residual graph G_f and has to check whether the edge is still in M for the next search.

There are at most n phases. Hence, total cost is $\mathcal{O}(mn^2)$.

How to choose augmenting paths?

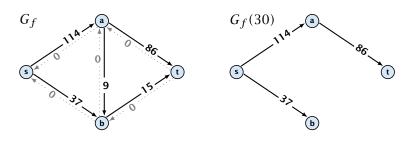
- We need to find paths efficiently.
- We want to guarantee a small number of iterations.

Several possibilities:

- Choose path with maximum bottleneck capacity.
- Choose path with sufficiently large bottleneck capacity.
- Choose the shortest augmenting path.

Intuition:

- Choosing a path with the highest bottleneck increases the flow as much as possible in a single step.
- Don't worry about finding the exact bottleneck.
- ▶ Maintain scaling parameter Δ .
- ▶ $G_f(\Delta)$ is a sub-graph of the residual graph G_f that contains only edges with capacity at least Δ .



```
Algorithm 1 maxflow(G, s, t, c)
 1: foreach e \in E do f_e \leftarrow 0;
 2: \Delta \leftarrow 2^{\lceil \log_2 C \rceil}
 3: while \Delta \geq 1 do
 4: G_f(\Delta) \leftarrow \Delta-residual graph
5: while there is augmenting path P in G_f(\Delta) do
6: f \leftarrow \text{augment}(f, c, P)
7: \text{update}(G_f(\Delta))
8: \Delta \leftarrow \Delta/2
  9: return f
```

Assumption:

All capacities are integers between 1 and C.

Invariant:

All flows and capacities are/remain integral throughout the algorithm.

Correctness:

The algorithm computes a maxflow:

- because of integrality we have $G_f(1) = G_f$
- therefore after the last phase there are no augmenting paths anymore
- this means we have a maximum flow.

Lemma 59

There are $\lceil \log C \rceil + 1$ iterations over Δ .

Proof: obvious.

Lemma 60

Let f be the flow at the end of a Δ -phase. Then the maximum flow is smaller than $val(f) + m\Delta$.

Proof: less obvious, but simple:

- ▶ There must exist an s-t cut in $G_f(\Delta)$ of zero capacity.
- ▶ In G_f this cut can have capacity at most $m\Delta$.
- This gives me an upper bound on the flow that I can still add.

Lemma 61

There are at most 2m augmentations per scaling-phase.

Proof:

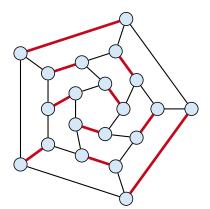
- Let f be the flow at the end of the previous phase.
- $ightharpoonup \operatorname{val}(f^*) \le \operatorname{val}(f) + 2m\Delta$
- **Each** augmentation increases flow by Δ .

Theorem 62

We need $\mathcal{O}(m \log C)$ augmentations. The algorithm can be implemented in time $\mathcal{O}(m^2 \log C)$.

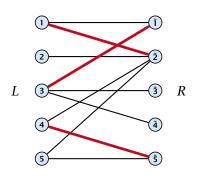
Matching

- ▶ Input: undirected graph G = (V, E).
- ► $M \subseteq E$ is a matching if each node appears in at most one edge in M.
- Maximum Matching: find a matching of maximum cardinality



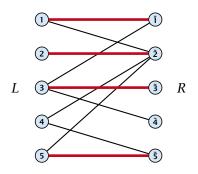
Bipartite Matching

- ▶ Input: undirected, bipartite graph $G = (L \uplus R, E)$.
- ▶ $M \subseteq E$ is a matching if each node appears in at most one edge in M.
- Maximum Matching: find a matching of maximum cardinality



Bipartite Matching

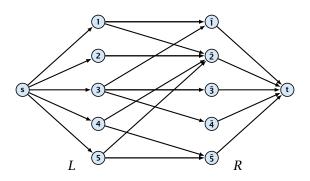
- ▶ Input: undirected, bipartite graph $G = (L \uplus R, E)$.
- ▶ $M \subseteq E$ is a matching if each node appears in at most one edge in M.
- Maximum Matching: find a matching of maximum cardinality



12.1 Matching

Maxflow Formulation

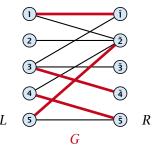
- ▶ Input: undirected, bipartite graph $G = (L \uplus R \uplus \{s, t\}, E')$.
- ▶ Direct all edges from *L* to *R*.
- Add source s and connect it to all nodes on the left.
- Add t and connect all nodes on the right to t.
- All edges have unit capacity.

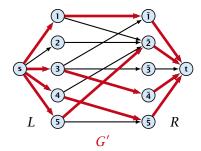


Proof

Max cardinality matching in $G \le \text{value of maxflow in } G'$

- Given a maximum matching M of cardinality k.
- Consider flow f that sends one unit along each of k paths.
- ightharpoonup f is a flow and has cardinality k.

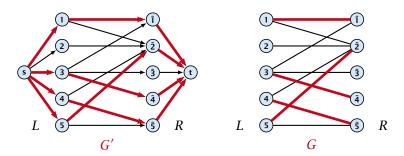




Proof

Max cardinality matching in $G \ge \text{value of maxflow in } G'$

- Let f be a maxflow in G' of value k
- ▶ Integrality theorem $\Rightarrow k$ integral; we can assume f is 0/1.
- Consider M= set of edges from L to R with f(e) = 1.
- ▶ Each node in L and R participates in at most one edge in M.
- |M| = k, as the flow must use at least k middle edges.



12. Jan. 2025 462/530

12.1 Matching

Which flow algorithm to use?

- Generic augmenting path: $\mathcal{O}(m \operatorname{val}(f^*)) = \mathcal{O}(mn)$.
- Capacity scaling: $\mathcal{O}(m^2 \log C) = \mathcal{O}(m^2)$.
- Shortest augmenting path: $O(mn^2)$.

For unit capacity simple graphs shortest augmenting path can be implemented in time $\mathcal{O}(m\sqrt{n})$.

A graph is a unit capacity simple graph if

- every edge has capacity 1
- a node has either at most one leaving edge or at most one entering edge



Baseball Elimination

team	wins	losses	remaining games			
i	w_i	ℓ_i	Atl	Phi	NY	Mon
Atlanta	83	71	_	1	6	1
Philadelphia	80	79	1	-	0	2
New York	78	78	6	0	_	0
Montreal	77	82	1	2	0	_

Which team can end the season with most wins?

- Montreal is eliminated, since even after winning all remaining games there are only 80 wins.
- But also Philadelphia is eliminated. Why?



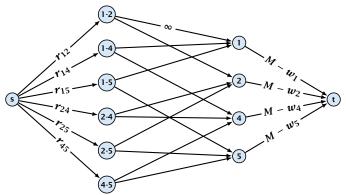
Baseball Elimination

Formal definition of the problem:

- ▶ Given a set S of teams, and one specific team $z \in S$.
- ▶ Team x has already won w_x games.
- ► Team x still has to play team y, r_{xy} times.
- Does team z still have a chance to finish with the most number of wins.

Baseball Elimination

Flow network for z = 3. M is number of wins Team 3 can still obtain.



Idea. Distribute the results of remaining games in such a way that no team gets too many wins.

Certificate of Elimination

Let $T \subseteq S$ be a subset of teams. Define

$$w(T) := \sum_{i \in T} w_i, \qquad r(T) := \sum_{i,j \in T, i < j} r_{ij}$$
 wins of teams in T remaining games among teams in T

If $\frac{w(T)+r(T)}{|T|}>M$ then one of the teams in T will have more than M wins in the end. A team that can win at most M games is therefore eliminated.

Theorem 63

A team z is eliminated if and only if the flow network for z does not allow a flow of value $\sum_{ij \in S \setminus \{z\}, i < j} \gamma_{ij}$.

Proof (⇐)

- Consider the mincut A in the flow network. Let T be the set of team-nodes in A.
- If for node x-y not both team-nodes x and y are in T, then x- $y \notin A$ as otw. the cut would cut an infinite capacity edge.
- We don't find a flow that saturates all source edges:

$$r(S \setminus \{z\}) > \operatorname{cap}(A, V \setminus A)$$

$$\geq \sum_{i < j: i \notin T \lor j \notin T} r_{ij} + \sum_{i \in T} (M - w_i)$$

$$\geq r(S \setminus \{z\}) - r(T) + |T|M - w(T)$$

▶ This gives M < (w(T) + r(T))/|T|, i.e., z is eliminated.

Baseball Elimination

Proof (⇒)

- Suppose we have a flow that saturates all source edges.
- We can assume that this flow is integral.
- For every pairing x-y it defines how many games team x and team y should win.
- The flow leaving the team-node x can be interpreted as the additional number of wins that team x will obtain.
- ▶ This is less than $M w_x$ because of capacity constraints.
- Hence, we found a set of results for the remaining games, such that no team obtains more than M wins in total.
- Hence, team z is not eliminated.



Project Selection

Project selection problem:

- Set P of possible projects. Project v has an associated profit p_v (can be positive or negative).
- Some projects have requirements (taking course EA2 requires course EA1).
- Dependencies are modelled in a graph. Edge (u, v) means "can't do project u without also doing project v."
- ▶ A subset *A* of projects is **feasible** if the prerequisites of every project in *A* also belong to *A*.

Goal: Find a feasible set of projects that maximizes the profit.

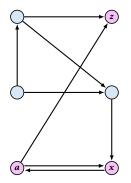


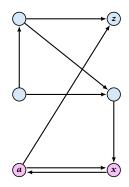
470/530

Project Selection

The prerequisite graph:

- \blacktriangleright {x, a, z} is a feasible subset.
- \triangleright {x, a} is infeasible.

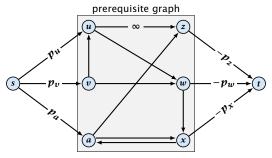




Project Selection

Mincut formulation:

- Edges in the prerequisite graph get infinite capacity.
- Add edge (s, v) with capacity p_v for nodes v with positive profit.
- Create edge (v,t) with capacity $-p_v$ for nodes v with negative profit.



Theorem 64

A is a mincut if $A \setminus \{s\}$ is the optimal set of projects.

Proof.

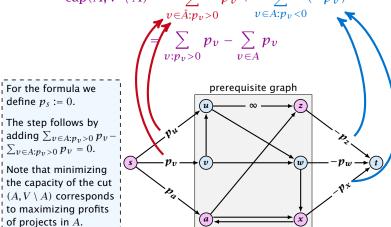
For the formula we

 $\sum_{v \in A: p_v > 0} p_v = 0.$

of projects in A.

define $p_s := 0$.

- A is feasible because of capacity infinity edges.
- $ightharpoonup \operatorname{cap}(A, V \setminus A) = \sum_{v \in \mathcal{V}} p_v + \sum_{v \in \mathcal{V}} (-p_v)$



Definition 65

An (s,t)-preflow is a function $f:E\mapsto \mathbb{R}^+$ that satisfies

1. For each edge e

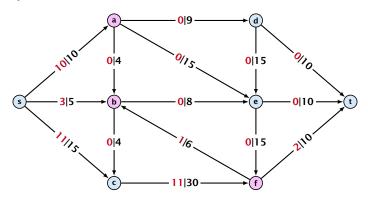
$$0 \le f(e) \le c(e)$$
.

(capacity constraints)

2. For each $v \in V \setminus \{s, t\}$

$$\sum_{e \in \text{out}(v)} f(e) \le \sum_{e \in \text{into}(v)} f(e) .$$

Example 66



A node that has $\sum_{e \in \text{out}(v)} f(e) < \sum_{e \in \text{into}(v)} f(e)$ is called an active node.

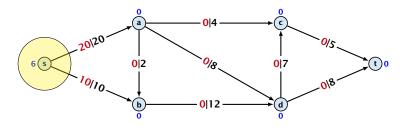
Definition:

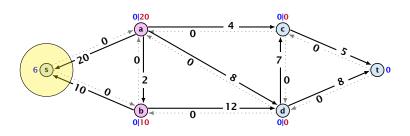
A labelling is a function $\ell: V \to \mathbb{N}$. It is valid for preflow f if

- $\ell(u) \le \ell(v) + 1$ for all edges (u, v) in the residual graph G_f (only non-zero capacity edges!!!)
- \blacktriangleright $\ell(s) = n$
- $\blacktriangleright \ell(t) = 0$

Intuition:

The labelling can be viewed as a height function. Whenever the height from node u to node v decreases by more than 1 (i.e., it goes very steep downhill from u to v), the corresponding edge must be saturated.





Lemma 67

A preflow that has a valid labelling saturates a cut.

Proof:

- ▶ There are n nodes but n + 1 different labels from 0, ..., n.
- ▶ There must exist a label $d \in \{0, ..., n\}$ such that none of the nodes carries this label.
- ▶ Let $A = \{v \in V \mid \ell(v) > d\}$ and $B = \{v \in V \mid \ell(v) < d\}$.
- ▶ We have $s \in A$ and $t \in B$ and there is no edge from A to B in the residual graph G_f ; this means that (A,B) is a saturated cut.

Lemma 68

A flow that has a valid labelling is a maximum flow.



Push Relabel Algorithms

Idea:

- start with some preflow and some valid labelling
- successively change the preflow while maintaining a valid labelling
- stop when you have a flow (i.e., no more active nodes)

Note that this is somewhat dual to an augmenting path algorithm. The former maintains the property that it has a feasible flow. It successively changes this flow until it saturates some cut in which case we conclude that the flow is maximum. A preflow push algorithm maintains the property that it has a saturated cut. The preflow is changed iteratively until it fulfills conservation constraints in which case we can conclude that we have a maximum flow.

Changing a Preflow

An arc (u, v) with $c_f(u, v) > 0$ in the residual graph is admissible if $\ell(u) = \ell(v) + 1$ (i.e., it goes downwards w.r.t. labelling ℓ).

The push operation

Consider an active node u with excess flow

$$f(u) = \sum_{e \in \text{into}(u)} f(e) - \sum_{e \in \text{out}(u)} f(e)$$
 and suppose $e = (u, v)$ is an admissible arc with residual capacity $c_f(e)$.

We can send flow $\min\{c_f(e), f(u)\}$ along e and obtain a new preflow. The old labelling is still valid (!!!).

- ▶ saturating push: $min\{f(u), c_f(e)\} = c_f(e)$ the arc e is deleted from the residual graph
- ▶ deactivating push: $min{f(u), c_f(e)} = f(u)$ the node u becomes inactive

Note that a push-operation may be saturating **and** deactivating at the same time.

Push Relabel Algorithms

The relabel operation

Consider an active node u that does not have an outgoing admissible arc.

Increasing the label of u by 1 results in a valid labelling.

- ▶ Edges (w, u) incoming to u still fulfill their constraint $\ell(w) \le \ell(u) + 1$.
- An outgoing edge (u, w) had $\ell(u) < \ell(w) + 1$ before since it was not admissible. Now: $\ell(u) \le \ell(w) + 1$.

Push Relabel Algorithms

Intuition:

We want to send flow downwards, since the source has a height/label of n and the target a height/label of 0. If we see an active node u with an admissible arc we push the flow at u towards the other end-point that has a lower height/label. If we do not have an admissible arc but excess flow into u it should roughly mean that the level/height/label of u should rise. (If we consider the flow to be water then this would be natural.)

Note that the above intuition is very incorrect as the labels are integral, i.e., they cannot really be seen as the height of a node.

Reminder

- In a preflow nodes may not fulfill conservation constraints; a node may have more incoming flow than outgoing flow.
- Such a node is called active.
- ▶ A labelling is valid if for every edge (u, v) in the residual graph $\ell(u) \le \ell(v) + 1$.
- An arc (u, v) in residual graph is admissible if $\ell(u) = \ell(v) + 1$.
- A saturating push along e pushes an amount of c(e) flow along the edge, thereby saturating the edge (and making it dissappear from the residual graph).
- A deactivating push along e = (u, v) pushes a flow of f(u), where f(u) is the excess flow of u. This makes u inactive.



Push Relabel Algorithms

```
Algorithm 1 maxflow(G, s, t, c)

1: find initial preflow f

2: while there is active node u do

3: if there is admiss. arc e out of u then

4: push(G, e, f, c)

5: else

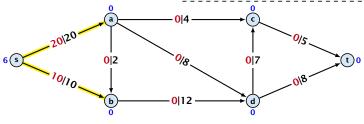
6: relabel(u)

7: return f
```

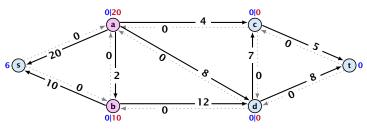
In the following example we always stick to the same active node \boldsymbol{u} until it becomes inactive but this is not required.

Preflow Push

The yellow edges indicate the cut that is introduced by the smallest missing label.



d**satetiathirilggpgis** hh



Note that the lemma is almost trivial. A node v having excess flow means that the current preflow ships something to v. The residual graph allows to undo flow. Therefore, there must exist a path that can undo the shipment and move it back to s. However, a formal proof is required.

Lemma 69

An active node has a path to s in the residual graph.

Proof.

- Let A denote the set of nodes that can reach s, and let B denote the remaining nodes. Note that $s \in A$.
- In the following we show that a node $b \in B$ has excess flow f(b) = 0 which gives the lemma.
- In the residual graph there are no edges into A, and, hence, no edges leaving A/entering B can carry any flow.
- Let $f(B) = \sum_{v \in B} f(v)$ be the excess flow of all nodes in B.

486/530

Let $f: E \to \mathbb{R}_0^+$ be a preflow. We introduce the notation

$$f(x,y) = \begin{cases} 0 & (x,y) \notin E \\ f((x,y)) & (x,y) \in E \end{cases}$$

We have

$$\begin{split} f(B) &= \sum_{b \in B} f(b) \\ &= \sum_{b \in B} \left(\sum_{v \in V} f(v, b) - \sum_{v \in V} f(b, v) \right) \\ &= \sum_{b \in B} \left(\sum_{v \in A} f(v, b) + \sum_{v \in B} f(v, b) - \sum_{v \in A} f(b, v) - \sum_{v \in B} f(b, v) \right) \\ &= -\sum_{b \in B} \sum_{v \in A} f(b, v) \\ &\leq 0 \end{split}$$

Hence, the excess flow f(b) must be 0 for every node $b \in B$.

Lemma 70

The label of a node cannot become larger than 2n-1.

Proof.

When increasing the label at a node u there exists a path from u to s of length at most n-1. Along each edge of the path the height/label can at most drop by 1, and the label of the source is n.

Lemma 71

There are only $O(n^2)$ relabel operations.

Lemma 72

The number of saturating pushes performed is at most O(mn).

Proof.

- Suppose that we just made a saturating push along (u, v).
- ▶ Hence, the edge (u, v) is deleted from the residual graph.
- For the edge to appear again, a push from v to u is required.
- Currently, $\ell(u) = \ell(v) + 1$, as we only make pushes along admissible edges.
- For a push from v to u the edge (v, u) must become admissible. The label of v must increase by at least 2.
- Since the label of v is at most 2n-1, there are at most n pushes along (u,v).

Lemma 73

The number of deactivating pushes performed is at most $O(n^2m)$.

Proof.

- ▶ Define a potential function $\Phi(f) = \sum_{\text{active nodes } v} \ell(v)$
- A saturating push increases Φ by $\leq 2n$ (when the target node becomes active it may contribute at most 2n to the sum).
- ▶ A relabel increases Φ by at most 1.
- A deactivating push decreases Φ by at least 1 as the node that is pushed from becomes inactive and has a label that is strictly larger than the target.
- Hence,

```
#deactivating_pushes \leq #relabels + 2n \cdot #saturating_pushes \leq \mathcal{O}(n^2m).
```

Theorem 74

There is an implementation of the generic push relabel algorithm with running time $\mathcal{O}(n^2m)$.

Proof:

For every node maintain a list of admissible edges starting at that node. Further maintain a list of active nodes.

A push along an edge (u, v) can be performed in constant time

- lacksquare check whether edge (v,u) needs to be added to G_f
- check whether (u, v) needs to be deleted (saturating push)
- check whether u becomes inactive and has to be deleted from the set of active nodes

A relabel at a node u can be performed in time O(n)

- check for all outgoing edges if they become admissible
- check for all incoming edges if they become non-admissible



For special variants of push relabel algorithms we organize the neighbours of a node into a linked list (possible neighbours in the residual graph G_f). Then we use the discharge-operation:

```
Algorithm 2 discharge(u)
 1: while u is active do
        v \leftarrow u.current-neighbour
 2:
      if v = \text{null then}
3:
              relabel(u)
4:
 5:
              u.current-neighbour ← u.neighbour-list-head
        else
6:
 7:
              if (u, v) admissible then push(u, v)
              else u.current-neighbour \leftarrow v.next-in-list
 8:
```

Note that *u.current-neighbour* is a global variable. It is only changed within the discharge routine, but keeps its value between consecutive calls to discharge.

Lemma 75

If v = null in Line 3, then there is no outgoing admissible edge from u.

In order for e to become admissible the other end-point say v has to push flow to u (so that the edge (u,v) re-appears in the residual graph). For this the label of v needs to be larger than the label of u. Then in order to make (u,v) admissible the label of u has to increase.

Proof.

- ▶ While pushing from *u* the current-neighbour pointer is only advanced if the current edge is not admissible.
- The only thing that could make the edge admissible again would be a relabel at u.
- If we reach the end of the list (v = null) all edges are not admissible.

This shows that discharge(u) is correct, and that we can perform a relabel in Line 4.



```
Algorithm 1 relabel-to-front(G, s, t)
1: initialize preflow
2: initialize node list L containing V \setminus \{s, t\} in any order
3: foreach u \in V \setminus \{s, t\} do
         u.current-neighbour \leftarrow u.neighbour-list-head
4:
5: u \leftarrow L.head
6: while u \neq \text{null do}
         old-height \leftarrow \ell(u)
7:
         discharge(u)
8:
         if \ell(u) > old-height then // relabel happened
9:
10:
               move u to the front of L
11:
         u \leftarrow u.next
```

Lemma 76 (Invariant)

In Line 6 of the relabel-to-front algorithm the following invariant holds.

- 1. The sequence L is topologically sorted w.r.t. the set of admissible edges; this means for an admissible edge (x,y) the node x appears before y in sequence L.
- **2.** No node before u in the list L is active.

Proof:

- Initialization:
 - 1. In the beginning s has label $n \ge 2$, and all other nodes have label 0. Hence, no edge is admissible, which means that any ordering L is permitted.
 - 2. We start with u being the head of the list; hence no node before u can be active

Maintenance:

- Pushes do no create any new admissible edges. Therefore, if discharge() does not relabel u, L is still topologically sorted.
 - After relabeling, u cannot have admissible incoming edges as such an edge (x, u) would have had a difference $\ell(x) \ell(u) \ge 2$ before the re-labeling (such edges do not exist in the residual graph).

Hence, moving u to the front does not violate the sorting property for any edge; however it fixes this property for all admissible edges leaving u that were generated by the relabeling.

Proof:

- Maintenance:
 - 2. If we do a relabel there is nothing to prove because the only node before u' (u in the next iteration) will be the current u; the discharge(u) operation only terminates when u is not active anymore.

For the case that we do not relabel, observe that the only way a predecessor could be active is that we push flow to it via an admissible arc. However, all admissible arc point to successors of u.

Note that the invariant means that for u = null we have a preflow with a valid labelling that does not have active nodes. This means we have a maximum flow.

Lemma 77

There are at most $O(n^3)$ calls to discharge(u).

Every discharge operation without a relabel advances u (the current node within list L). Hence, if we have n discharge operations without a relabel we have $u = \mathrm{null}$ and the algorithm terminates.

Therefore, the number of calls to discharge is at most $n(\#relabels + 1) = \mathcal{O}(n^3)$.

Lemma 78

The cost for all relabel-operations is only $\mathcal{O}(n^2)$.

A relabel-operation at a node is constant time (increasing the label and resetting u.current-neighbour). In total we have $\mathcal{O}(n^2)$ relabel-operations.

Recall that a saturating push operation $(\min\{c_f(e), f(u)\} = c_f(e))$ can also be a deactivating push operation $(\min\{c_f(e), f(u)\} = f(u))$.

Lemma 79

The cost for all saturating push-operations that are **not** deactivating is only O(mn).

Note that such a push-operation leaves the node u active but makes the edge e disappear from the residual graph. Therefore the push-operation is immediately followed by an increase of the pointer u.current-neighbour.

This pointer can traverse the neighbour-list at most $\mathcal{O}(n)$ times (upper bound on number of relabels) and the neighbour-list has only degree(u) + 1 many entries (+1 for null-entry).

Lemma 80

The cost for all deactivating push-operations is only $O(n^3)$.

A deactivating push-operation takes constant time and ends the current call to discharge(). Hence, there are only $\mathcal{O}(n^3)$ such operations.

Theorem 81

The push-relabel algorithm with the rule relabel-to-front takes time $\mathcal{O}(n^3)$.

13.3 Highest Label

Algorithm 1 highest-label (G, s, t)

- 1: initialize preflow
- 2: foreach $u \in V \setminus \{s, t\}$ do
- 3: $u.current-neighbour \leftarrow u.neighbour-list-head$
- 4: **while** \exists active node u **do**
- 5: select active node u with highest label
- 6: discharge(u)

13.3 Highest Label

Lemma 82

When using highest label the number of deactivating pushes is only $\mathcal{O}(n^3)$.

A push from a node on level ℓ can only "activate" nodes on levels strictly less than ℓ .

This means, after a deactivating push from \boldsymbol{u} a relabel is required to make \boldsymbol{u} active again.

Hence, after n deactivating pushes without an intermediate relabel there are no active nodes left.

Therefore, the number of deactivating pushes is at most $n(\#relabels + 1) = O(n^3)$.

Since a discharge-operation is terminated by a deactivating push this gives an upper bound of $\mathcal{O}(n^3)$ on the number of discharge-operations.

The cost for relabels and saturating pushes can be estimated in exactly the same way as in the case of the generic push-relabel algorithm.

Question:

How do we find the next node for a discharge operation?

Maintain lists L_i , $i \in \{0, ..., 2n\}$, where list L_i contains active nodes with label i (maintaining these lists induces only constant additional cost for every push-operation and for every relabel-operation).

After a discharge operation terminated for a node u with label k, traverse the lists $L_k, L_{k-1}, \ldots, L_0$, (in that order) until you find a non-empty list.

Unless the last (deactivating) push was to s or t the list k-1 must be non-empty (i.e., the search takes constant time).

Hence, the total time required for searching for active nodes is at most

$$O(n^3) + n(\# deactivating-pushes-to-s-or-t)$$

Lemma 83

The number of deactivating pushes to s or t is at most $O(n^2)$.

With this lemma we get

Theorem 84

The push-relabel algorithm with the rule highest-label takes time $\mathcal{O}(n^3)$.

Proof of the Lemma.

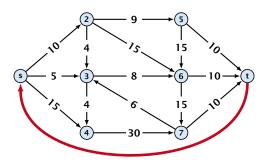
- We only show that the number of pushes to the source is at most $\mathcal{O}(n^2)$. A similar argument holds for the target.
- After a node v (which must have $\ell(v) = n+1$) made a deactivating push to the source there needs to be another node whose label is increased from $\leq n+1$ to n+2 before v can become active again.
- This happens for every push that v makes to the source. Since, every node can pass the threshold n + 2 at most once, v can make at most n pushes to the source.
- As this holds for every node the total number of pushes to the source is at most $\mathcal{O}(n^2)$.

Problem Definition:

```
\begin{aligned} & \min & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E : & 0 \leq f(e) \leq u(e) \\ & & \forall v \in V : & f(v) = b(v) \end{aligned}
```

- G = (V, E) is a directed graph.
- ▶ $u: E \to \mathbb{R}_0^+ \cup \{\infty\}$ is the capacity function.
- ► $c: E \to \mathbb{R}$ is the cost function (note that c(e) may be negative).
- ▶ $b: V \to \mathbb{R}$, $\sum_{v \in V} b(v) = 0$ is a demand function.

Solve Maxflow Using Mincost Flow



- Given a flow network for a standard maxflow problem.
- Set b(v) = 0 for every node. Keep the capacity function u for all edges. Set the cost c(e) for every edge to 0.
- ▶ Add an edge from t to s with infinite capacity and cost -1.
- ► Then, $val(f^*) = -cost(f_{min})$, where f^* is a maxflow, and f_{min} is a mincost-flow.



Solve Maxflow Using Mincost Flow

Solve decision version of maxflow:

- Given a flow network for a standard maxflow problem, and a value k.
- Set b(v) = 0 for every node apart from s or t. Set b(s) = -k and b(t) = k.
- Set edge-costs to zero, and keep the capacities.
- There exists a maxflow of value at least k if and only if the mincost-flow problem is feasible.

Generalization

Our model:

min
$$\sum_{e} c(e) f(e)$$

s.t. $\forall e \in E: 0 \le f(e) \le u(e)$
 $\forall v \in V: f(v) = b(v)$

where
$$b: V \to \mathbb{R}$$
, $\sum_{v} b(v) = 0$; $u: E \to \mathbb{R}_0^+ \cup \{\infty\}$; $c: E \to \mathbb{R}$;

A more general model?

$$\begin{aligned} & \text{min} & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & \ell(e) \leq f(e) \leq u(e) \\ & & & \forall v \in V: & a(v) \leq f(v) \leq b(v) \end{aligned}$$

where $a: V \to \mathbb{R}$, $b: V \to \mathbb{R}$; $\ell: E \to \mathbb{R} \cup \{-\infty\}$, $u: E \to \mathbb{R} \cup \{\infty\}$ $c: E \to \mathbb{R}$;

Generalization

Differences

- Flow along an edge e may have non-zero lower bound $\ell(e)$.
- Flow along e may have negative upper bound u(e).
- ▶ The demand at a node v may have lower bound a(v) and upper bound b(v) instead of just lower bound = upper bound = b(v).

12. Jan. 2025

Reduction I

$$\begin{aligned} & \min \quad \sum_{e} c(e) f(e) \\ & \text{s.t.} \quad \forall e \in E: \ \ell(e) \leq f(e) \leq u(e) \\ & \quad \forall v \in V: \ a(v) \leq f(v) \leq b(v) \end{aligned}$$

We can assume that a(v) = b(v):

Add new node r.

Add edge (r, v) for all $v \in V$.

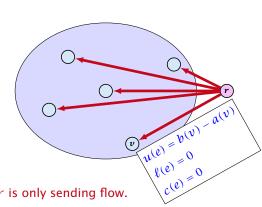
Set $\ell(e) = c(e) = 0$ for these edges.

Set u(e) = b(v) - a(v) for edge (r, v).

Set a(v) = b(v) for all $v \in V$.

Set $b(r) = -\sum_{v \in V} b(v)$.

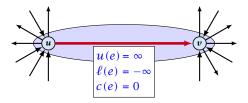
 $-\sum_{v}b(v)$ is negative; hence r is only sending flow.



Reduction II

$$\begin{aligned} & \text{min} & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E : & \ell(e) \leq f(e) \leq u(e) \\ & & \forall v \in V : & f(v) = b(v) \end{aligned}$$

We can assume that either $\ell(e) \neq -\infty$ or $u(e) \neq \infty$:

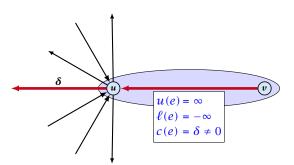


If c(e) = 0 we can contract the edge/identify nodes u and v.

If $c(e) \neq 0$ we can transform the graph so that c(e) = 0.

Reduction II

We can transform any network so that a particular edge has c(e) = 0:

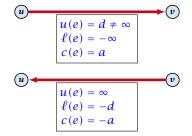


Additionally we set b(u) = 0.

Reduction III

$$\begin{aligned} & \min & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E : & \ell(e) \leq f(e) \leq u(e) \\ & & \forall v \in V : & f(v) = b(v) \end{aligned}$$

We can assume that $\ell(e) \neq -\infty$:

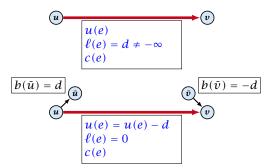


Replace the edge by an edge in opposite direction.

Reduction IV

$$\begin{aligned} & \min & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & \ell(e) \leq f(e) \leq u(e) \\ & & & \forall v \in V: & f(v) = b(v) \end{aligned}$$

We can assume that $\ell(e) = 0$:



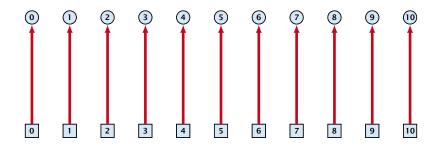
The added edges have infinite capacity and cost c(e)/2.

Applications

Caterer Problem

- She needs to supply r_i napkins on N successive days.
- She can buy new napkins at p cents each.
- She can launder them at a fast laundry that takes m days and cost f cents a napkin.
- She can use a slow laundry that takes k > m days and costs s cents each.
- At the end of each day she should determine how many to send to each laundry and how many to buy in order to fulfill demand.
- Minimize cost.



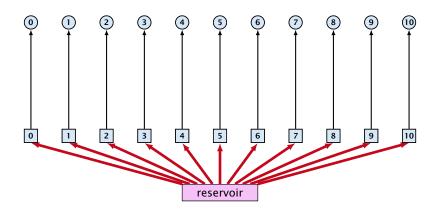


day edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = r_i$;

cost: c(e) = 0

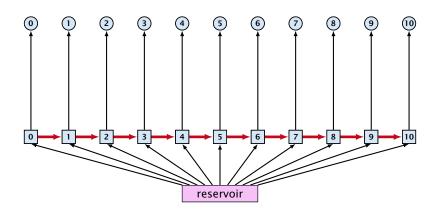


buy edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = 0$;

cost: c(e) = p

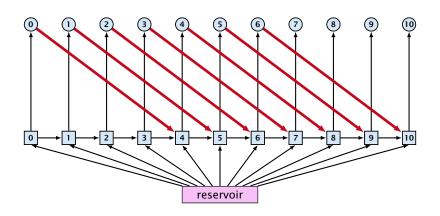


forward edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = 0$;

cost: c(e) = 0

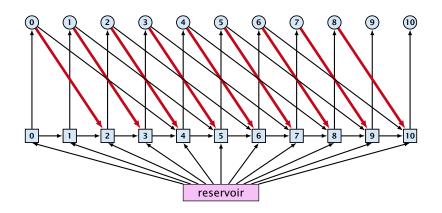


slow edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = 0$;

cost: c(e) = s

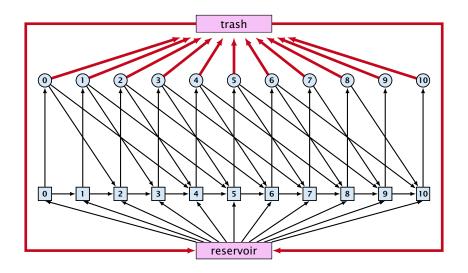


fast edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = 0$;

cost: c(e) = f



trash edges:

upper bound: $u(e_i) = \infty$; lower bound: $\ell(e_i) = 0$;

cost: c(e) = 0

Residual Graph

Version A:

The residual graph G' for a mincost flow is just a copy of the graph G.

If we send f(e) along an edge, the corresponding edge e' in the residual graph has its lower and upper bound changed to $\ell(e') = \ell(e) - f(e)$ and u(e') = u(e) - f(e).

Version B:

The residual graph for a mincost flow is exactly defined as the residual graph for standard flows, with the only exception that one needs to define a cost for the residual edge.

For a flow of z from u to v the residual edge (v,u) has capacity z and a cost of -c((u,v)).



A circulation in a graph G = (V, E) is a function $f : E \to \mathbb{R}^+$ that has an excess flow f(v) = 0 for every node $v \in V$.

A circulation is feasible if it fulfills capacity constraints, i.e., $f(e) \le u(e)$ for every edge of G.

Lemma 85

A given flow is a mincost-flow if and only if the corresponding residual graph G_f does not have a feasible circulation of negative cost.

 \Rightarrow Suppose that g is a feasible circulation of negative cost in the residual graph.

Then f+g is a feasible flow with cost $\cos(f)+\cos(g)<\cos(f)$. Hence, f is not minimum cost.

 \leftarrow Let f be a non-mincost flow, and let f^* be a min-cost flow. We need to show that the residual graph has a feasible circulation with negative cost.

Clearly $f^* - f$ is a circulation of negative cost. One can also easily see that it is feasible for the residual graph. (after sending -f in the residual graph (pushing all flow back) we arrive at the original graph; for this f^* is clearly feasible)

For previous slide:

 $g = f^* - f$ is obtained by computing $\Delta(e) = f^*(e) - f(e)$ for every edge e = (u, v). If the result is positive set $g((u, v)) = \Delta(e)$ and g((v, u)) = 0. Otherwise set g((u, v)) = 0 and $g((v, u)) = -\Delta(e)$.



Lemma 86

A graph (without zero-capacity edges) has a feasible circulation of negative cost if and only if it has a negative cycle w.r.t. edge-weights $c : E \to \mathbb{R}$.

Proof.

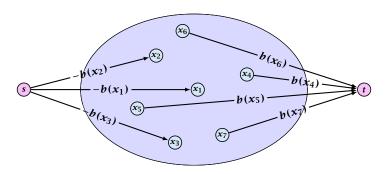
- Suppose that we have a negative cost circulation.
- Find directed cycle only using edges that have non-zero flow.
- If this cycle has negative cost you are done.
- Otherwise send flow in opposite direction along the cycle until the bottleneck edge(s) does not carry any flow.
- You still have a circulation with negative cost.
- Repeat.



Algorithm 48 CycleCanceling(G = (V, E), c, u, b)

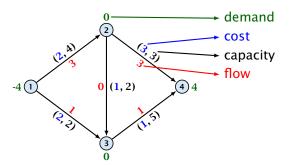
- 1: establish a feasible flow f in G
- 2: **while** G_f contains negative cycle **do**
- 3: use Bellman-Ford to find a negative circuit Z
- 4: $\delta \leftarrow \min\{u_f(e) \mid e \in Z\}$
- 5: augment δ units along Z and update G_f

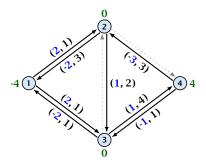
How do we find the initial feasible flow?

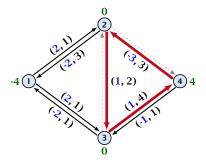


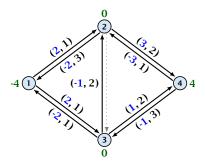
- Connect new node s to all nodes with negative b(v)-value.
- Connect nodes with positive b(v)-value to a new node t.
- ► There exist a feasible flow in the original graph iff in the resulting graph there exists an *s-t* flow of value

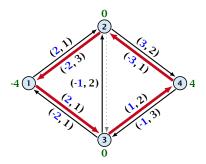
$$\sum_{v:b(v)<0} (-b(v)) = \sum_{v:b(v)>0} b(v) .$$

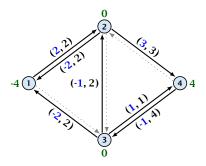












Lemma 87

The improving cycle algorithm runs in time $O(nm^2CU)$, for integer capacities and costs, when for all edges e, $|c(e)| \le C$ and $|u(e)| \le U$.

- Running time of Bellman-Ford is O(mn).
- ▶ Pushing flow along the cycle can be done in time O(n).
- Each iteration decreases the total cost by at least 1.
- ► The true optimum cost must lie in the interval [-mCU,...,+mCU].

Note that this lemma is weak since it does not allow for edges with infinite capacity.

A general mincost flow problem is of the following form:

$$\begin{aligned} & \min \quad \sum_{e} c(e) f(e) \\ & \text{s.t.} \quad \forall e \in E : \quad \ell(e) \leq f(e) \leq u(e) \\ & \quad \forall v \in V : \quad a(v) \leq f(v) \leq b(v) \end{aligned}$$
 where $a: V \to \mathbb{R}, \ b: V \to \mathbb{R}; \ \ell: E \to \mathbb{R} \cup \{-\infty\}, \ u: E \to \mathbb{R} \cup \{\infty\}$ $c: E \to \mathbb{R};$

Lemma 88 (without proof)

A general mincost flow problem can be solved in polynomial time.